SEQUENCE LISTING

Liu Zho Asu Ren Zha Zha Wan Weh	Tang, Y. T., Chenghua u, Ping ndi, Vinod , Feiyan o, Qing A. ng, Jie g, Jian-Rui rman, Tom anac, Radoj					
	Novel Nuclo	eic Acids a	nd			
<130>	789CIP2C					
<140> <141>	To Be Assig	gned				
<150>	09/574,454					
	2000-05-19					
<150>	09/519,705					
	2000-03-07					
<160>	35					
<170>	pt_FL_genes	s Version 2	. 0			
<210>	1					
<211>	_					
<212>	DNA					
<213>	Homo sapier	ns				
<220>						
<221>						
<222>	(258)(671	L)				
<400>						
taagcttgcg	gccgccccgg	ccgcccccgc	gggggacacc	teetegeege	cacagettta	60
ggaaacacct	ctacctcctg	cgaggcctcc	cgggctccgg	gaaaactaca	ctggccagac	120
aattgcagca	tgactttccc	agggccctga	ttttcagcac	ggatgatttt	ttcttcaggg	180

aagatggtgc ctatgagttc aatcctgact tcctggagga agctcatgaa tggaaccaaa	240
aaagagcaag aaaagca atg agg aat ggc ata tcc ccc att att att gat Met Arg Asn Gly Ile Ser Pro Ile Ile Asp 1 5 10	290
aat acc aac ctc cac gcc tgg gaa atg aag ccc tat gca gtc atg gca Asn Thr Asn Leu His Ala Trp Glu Met Lys Pro Tyr Ala Val Met Ala 15 20 25	338
ctt gaa aat aac tat gaa gtt ata ttc cga gaa cct gac act cgc tgg Leu Glu Asn Asn Tyr Glu Val Ile Phe Arg Glu Pro Asp Thr Arg Trp 30 35 40	386
aaa ttc aac gtt caa gag tta gca aga aga aac att cat ggt gtc tca Lys Phe Asn Val Gln Glu Leu Ala Arg Arg Asn Ile His Gly Val Ser 45 50 55	434
aga gaa aaa atc cac cga atg aaa gaa cgg tat gaa cac gat gtt act Arg Glu Lys Ile His Arg Met Lys Glu Arg Tyr Glu His Asp Val Thr 60 65 70 75	482
ttt cac agt gtg ctt cat gca gaa aag cca agc aga atg aac aga aac Phe His Ser Val Leu His Ala Glu Lys Pro Ser Arg Met Asn Arg Asn 80 85 90	530
cag gac agg aat aat gca ttg cct tcc aac aat gcc aga tac tgg aat Gln Asp Arg Asn Asn Ala Leu Pro Ser Asn Asn Ala Arg Tyr Trp Asn 95 100 105	578
tcc tac aca gag ttt cca aac cgg agg gcc cac ggt gga ttt aca aat Ser Tyr Thr Glu Phe Pro Asn Arg Arg Ala His Gly Gly Phe Thr Asn 110 115 120	626
gag agc tcc tat cac aga agg ggc ggt tgt cac cat gga tat tag agg Glu Ser Ser Tyr His Arg Arg Gly Gly Cys His His Gly Tyr * 125 130 135	674
cctatcttac agccaggcag aattttccta agtcagtttc tacttcagtt tttgttattt	734
tttgttgcat tttagtcaga gctccaattc cagtgtaaat agctgaactc aaaagtttct	794
gagcaaagtc attatattca ctttcttcac caaaatttgt taaagtgctt ctatatgcat	854
ggtctgatgc tgggaattct gcagatttga gtaaacagtc tctttctcta gggtaagaat	914
ttgaaaccaa aacttgagaa cacacccaag aatatattta cataggttca tagatgaaat	974
aaagtgttta tattatatat aagcttcagt accatttgct ctgaagtgat ctatttattt	1034
tttcaggaaa ttcatctcca tcggtaaagt tgggaaggtg gagagaagtg gtgggggggc	1094
attgctactt atcaaagtgc cattgctact ttgataatct atgtatctaa aaatgtgaga	1154
tgtgcgactc ttatgatact gattttcctt taatgttaat atgccagaaa gcatacatct	1214

aagggaacat tgtccttcaa agtagacact ttgggaagtt atttctttat tttaatgatg 1274 tatcattgtt aaaaatgctg tcaaatcctt aatagctaca ggagctactg agggaaatca 1334 gtgtcattat ttaaagtcac gccttgtgtt tttactactt tattcagcag gattaaacct 1394 gaataacttt tggctgttgt gctaatagtg taaataaaat aagcctgcct tcataaaaca 1454 ctaactttta aaaggaataa acgacttcta aaattatgcc tattaacatg tgtaattagt 1514 cggcagctca aatgtttggg agtgcaagaa attaggcacc ccaggatata ggtcatacag 1574 ggatatataa aagccatgct cattacaaaa tgagcagttg atgttttatg tggcattaag 1634 acaatcaagt cctcacaact ctggaatgtc ttcttatact gatgctgaat ttatgaatcc 1694 aaattaattt ccaacaggtt ggaatcagat ttaatgtgag atcatgatag acaagaccac 1754 agaggacgta tgctctattt cttgttggcc aacagcttct ttctaatgtt ctgtgaaaaa 1814 ttattttaag tgtcttatat aatggtgctt ttatggttat taaaaattgt aaatggtatc 1874 acatttatat ggatttgtca ttggatcttt ttttggttca acaataaaaa aatttaatta 1934 cctaaatgcc aagaaactca acaatatacc agtttttctg tatcacaggc ttatttacca 1994 gtctttttt taataaatag gaatcgtaaa ggtaatgaca aaagcagcct tataatttag 2054 ttgcttatat atttgatctg tgtacatgag actgttttaa cgttatctga cactactgaa 2114 acctgctcga catctccatg actaccaaca ccatgtgtaa tgttttcttc actaacattt 2174 taaaaactgg tatctccttt gagtaagttt ggctgacaat agtaaatccc aatgaatcta 2234 2235

<210> 2

<211> 4600

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (349)..(4305)

<400> 2

ttcaaggtca cgggccgtgc cagtccctac ccctagtgcc ccgcagcgtg ccagcccact 60 gatgccggca ggccgttacg ctatgcctgt ggcagcctcc cacctgggag aggactggcg 120 gccacgaggg caacttcgtg ggagaggtgg tggcgggctc ggggttcaga ggctgaagct 180 gggtcgcagc tcgaccgggg cgctgatctg tggctgtccg ttccttgctg gagaatttgg 240

ccacaaagag ctgcc	aagat agctgggd	cca ggaagaaagc	gccgcagccc tgacc	cagac 300
gctgttgccg acccc	ggggc actctggd	ctg tcgaccaagc	ggctcaag atg tct Met Ser 1	
			cca gga ctg gag Pro Gly Leu Glu 15	_
			aac cag gag agc Asn Gln Glu Ser	
			tat gtt gcc cag of Tyr Val Ala Gln .	
			gcc tca gcc tcc 6 Ala Ser Ala Ser 1 65	
	Thr Gly Ser Ar		cgg ctg ttc ttt (Arg Leu Phe Phe 1 80	
			gag cag acc aaa g Glu Gln Thr Lys (95	
			act ccc act cca of Thr Pro Thr Pro	
Leu Leu Leu Asp 7			gtg att gtc aag o Val Ile Val Lys 1 130	
		_	gta gat gct gct t Val Asp Ala Ala 1 145	
		g Phe Ala Gly	ggt cag cag tgg g Gly Gln Gln Trp (160	
			gct aaa gtg caa a Ala Lys Val Gln 1 175	
			aaa aag gtg cct a Lys Lys Val Pro N	
Leu Thr Trp Pro S			gaa cag ctt tgc a Glu Gln Leu Cys 1 210	

cca Pro	ccg Pro	ctg Leu	aac Asn 215	tcc Ser	caa Gln	acc Thr	tgc Cys	ctc Leu 220	ctg Leu	ggc Gly	tca Ser	gag Glu	gag Glu 225	aat Asn	tta Leu	1029
gcc Ala	cct Pro	ttg Leu 230	gca Ala	gga Gly	gag Glu	aaa Lys	gca Ala 235	gtg Val	cct Pro	ccc Pro	Gly ggg	aat Asn 240	gac Asp	cca Pro	gtc Val	1077
tct Ser	cca Pro 245	gcc Ala	atg Met	gtc Val	cgg Arg	agc Ser 250	aga Arg	aac Asn	cct Pro	ggg Gly	aaa Lys 255	gat Asp	gac Asp	tgt Cys	gcc Ala	1125
aag Lys 260	gag Glu	gag Glu	atg Met	gca Ala	gtg Val 265	gca Ala	gca Ala	gat Asp	gct Ala	gca Ala 270	acc Thr	ttg Leu	gtg Val	gat Asp	gag Glu 275	1173
ccc Pro	gag Glu	tcg Ser	atg Met	gtg Val 280	aac Asn	ctg Leu	gcg Ala	ttt Phe	gtc Val 285	aag Lys	aat Asn	gac Asp	tcg Ser	tat Tyr 290	gag Glu	1221
aag Lys	ggc Gly	ccg Pro	gat Asp 295	tca Ser	gtg Val	gtg Val	gtg Val	cac His 300	gtg Val	tac Tyr	gtg Val	aag Lys	gag Glu 305	atc Ile	tgc Cys	1269
agg Arg	gac Asp	acc Thr 310	Ser	aga Arg	gta Val	ctt Leu	ttc Phe 315	Arg	gag Glu	cag Gln	gac Asp	ttc Phe 320	acg Thr	ctc Leu	atc Ile	1317
ttc Phe	cag Gln 325	Thr	agg Arg	gat Asp	gga Gly	aac Asn 330	ttc Phe	ctg Leu	agg Arg	ctg Leu	cac His	PLU	ggc Gly	tgt Cys	Gly ggg	1365
ccc Pro 340	His	acc Thr	acc Thr	ttc Phe	cgt Arg 345	Trp	cag Glr	gtg Val	aag Lys	cto Leu 350	ı Arg	g aat g Asn	ctg Leu	att Ile	gag Glu 355	1413
cca Pro	gag Glu	g cag Glr	g tgo n Cys	acc Thr	Phe	tgt Cys	tto Phe	acg Thr	g gct Ala 365	a Ser	cgo Arg	ato g Ile	gac Asp	ato 116 370	tgc Cys	1461
ctt Leu	cgt Arg	aag J Lys	g agg s Arg 375	g Glr	g agt n Ser	caç Glr	cgo Arg	tgg Trp 380	O GI	g ggo	c cto / Let	g gaq ı Glı	g gcc 1 Ala 385	LLC	g gct o Ala	1509
gca Ala	ı cga	g Val	l Gly	ggt Gly	gca / Ala	a aag a Lys	g gti Va: 39:	I Ala	c gto a Va	g ccq l Pro	g aca	a gg r Gl 40	y PI	a aco	c cct r Pro	1557
cto Lei	g gat 1 Asj 40	o Se	a ace	c cca r Pro	a cca o Pro	a gga o Gly 410	/ G1	t gc	t cc a Pr	c ca o Hi	c cc s Pr 41	о пе	g aca	a gge r Gl	c cag y Gln	1605
gaq Gli 420	u Gl	g gc u Al	c cg a Ar	g gc	t gtg a Va 42	1 G1	g aa ı Ly	g ga s As	t aa p Ly	a tc s Se 43	т. г	g gc s Al	a cg a Ar	a tc g Se	t gag r Glu 435	1653

gac Asp	aca Thr	Gly	cta Leu	gac Asp 440	Ser	gtg Val	gca Ala	acc Thr	cgc Arg 445	Thr	ccc Pro	atg Met	gag Glu	cat His 450	gta Val	1701
acc Thr	cca Pro	aag Lys	cca Pro 455	Glu	aca Thr	cac His	ctg Leu	gcc Ala 460	Ser	ccc	aag Lys	cct Pro	aca Thr 465	Cys	atg Met	1749
gtg Val	cct Pro	ccc Pro 470	atg Met	ccc Pro	cac His	agc Ser	cca Pro 475	Val	agt Ser	gga Gly	gac Asp	agc Ser 480	gtg Val	gag Glu	gag Glu	1797
		Glu	gaa Glu													1845
gtc Val 500	Asn	tta Leu	ggc Gly	aac Asn	acc Thr 505	tgc Cys	ttc Phe	atg Met	aac Asn	agc Ser 510	gtc Val	att Ile	cag Gln	tct Ser	ctg Leu 515	1893
tcc Ser	aac Asn	act Thr	cgg Arg	gaa Glu 520	ctc Leu	cgg Arg	gac Asp	ttc Phe	ttc Phe 525	cat His	gac Asp	cgc Arg	tcc Ser	ttt Phe 530	gag Glu	1941
gct Ala	gag Glu	atc Ile	aac Asn 535	tac Tyr	aac Asn	aac Asn	cca Pro	cta Leu 540	gly aaa	act Thr	ggt Gly	gly aaa	cgt Arg 545	ctg Leu	gcc Ala	1989
att Ile	ggc Gly	ttt Phe 550	gcc Ala	gtg Val	ctg Leu	ctt Leu	cgg Arg 555	gcg Ala	ctg Leu	tgg Trp	aag Lys	ggc Gly 560	acc Thr	cac His	cat His	2037
gcc Ala	ttc Phe 565	cag Gln	cct Pro	tcc Ser	aag Lys	ttg Leu 570	aag Lys	gcc Ala	att Ile	gtg Val	gcg Ala 575	agt Ser	aag Lys	gcc Ala	agc Ser	2085
cag Gln 580	ttc Phe	aca Thr	ggc Gly	tat Tyr	gca Ala 585	cag Gln	cat His	gat Asp	gcc Ala	cag Gln 590	gag Glu	ttc Phe	atg Met	gct Ala	ttc Phe 595	2133
ctg Leu	ctg Leu	gat Asp	gly ggg	ctg Leu 600	cac His	gag Glu	gac Asp	ctg Leu	aat Asn 605	cgc Arg	att Ile	cag Gln	aac Asn	aag Lys 610	ccc Pro	2181
tac Tyr	aca Thr	gag Glu	acc Thr 615	gtg Val	gat Asp	tca Ser	gat Asp	999 Gly 620	cgg Arg	ccc Pro	gat Asp	gag Glu	gtg Val 625	gta Val	gct Ala	2229
gag Glu	gaa Glu	gca Ala 630	tgg Trp	cag Gln	cgg Arg	cac His	aag Lys 635	atg Met	agg Arg	aat Asn	gac Asp	tct Ser 640	ttc Phe	atc Ile	gtg Val	2277
gac Asp	cta Leu 645	ttt Phe	cag Gln	gly ggg	cag Gln	tac Tyr 650	aag Lys	tcg Ser	aag Lys	ctg Leu	gtg Val 655	tgc Cys	cct Pro	gtg Val	tgt Cys	2325
gcc	aag	gtc	tcc	atc	act	ttt	gac	ccg	ttt	ctt	tat	ctg	ccg	gtg	ccc	2373

	Lys	Val	Ser	Ile	Thr 665	Phe	Asp	Pro	Phe	Leu 670	Tyr	Leu	Pro	Val	Pro 675	
660 ttg Leu	cca Pro	caa Gln	aag Lys	Gln	aag	gtt Val	ctc Leu	cct Pro	Val	ttt	tat Tyr	ttt Phe	gcc Ala	cga Arg	gag	2421
								Leu					Lys	690 gag Glu		2469
tcc Ser	act Thr	gcg Ala	agc Ser	gaa Glu	gta Val	ttg Leu	Asp	700 tcc Ser	ctc Leu	tct Ser	cag Gln	Ser	705 gtt Val	cat His	gtg Val	2517
aag Lys	cct Pro	710 gag Glu	aac Asn	ctg Leu	cgt Arg	ttg Leu	715 gcg Ala	gag Glu	gta Val	att Ile	aag Lys	720 aat Asn	cgt Arg	ttt Phe	cat His	2565
cgt Arg	725 gtg Val	ttc Phe	cta Leu	ccc Pro	tcc Ser	730 cac His	tca Ser	ctg Leu	gac Asp	act Thr	735 gtg Val	tcc Ser	cca Pro	tct Ser	gat Asp	2613
740 acg Thr	ctc Leu	ctc Leu	tgc Cys	ttt Phe	745 gag Glu	ctg Leu	cta Leu	tcc Ser	tca Ser	750 gag Glu	ttg Leu	gct Ala	aag Lys	gag Glu	755 cgg Arg	2661
gta	gtg	gtg	cta	760 gag	gtg	caa	cag	cgc	765 ccc	cag	gtg	ccc	agc	770 gtc Val	ccc	2709
atc	tcc	aag	775 tgt	gca	gcc	tgc	cag	780 cgg	aag	caa	cag	tcg	785 gag	gat Asp	gaa	2757
aag	ctg	790 aag	cgc	tgt	acc	cgg	795 tgc	tac	cgt	gtg	ggc	800 tac	tgc	aac	cag	2805
ctc	805 tgc	cag	aaa	acc	cac	810 tgg	cct	gac	cac	aag	815 ggc	ctc	tgc	Asn	cct	2853
820 gag	aac	att	ggc	tac	825 ccc	ttc	ctg	gtc	agt	830 gta	cct	gcc	tca	Arg	835 ctc	2901
				840					845					Arg 850 tac		2949
Thr	Tyr	Ala	Arg 855	Leu	Ala	Gln	Leu	Leu 860	Glu	Gly	Tyr	Ala	Arg 865	Tyr ttg	Ser	2997
Val	Ser	Val 870	Phe	Gln	Pro	Pro	Phe 875	Gln	Pro	Gly	Arg	Met 880	Ala	Leu	Glu	3045
Ser	Gln	Ser	Pro	Gly	Cys	Thr	Thr	Leu	Leu	Ser	Thr	Gly	Ser	Leu	Glu	3043

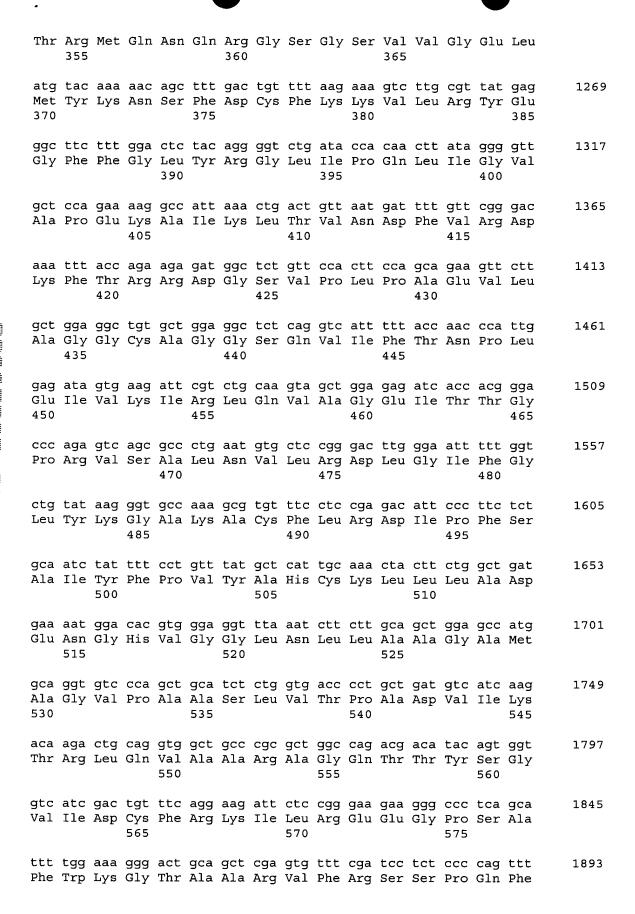
885 890 895 gct ggg gac agc gag aga gac ccc att cag cca cct gag ctc cag ctg 3093 Ala Gly Asp Ser Glu Arg Asp Pro Ile Gln Pro Pro Glu Leu Gln Leu 900 905 910 gtg acc cct atg gct gag ggg gac aca ggg ctt ccc cgg gtg tgg gca 3141 Val Thr Pro Met Ala Glu Gly Asp Thr Gly Leu Pro Arg Val Trp Ala 920 925 ged ect gad egg ggt ect gtg eec age ace agt gga att tet tet gag 3189 Ala Pro Asp Arg Gly Pro Val Pro Ser Thr Ser Gly Ile Ser Ser Glu 935 940 atg ctg gcc agt ggg ccc att gag gtt ggc tcc ttg cca gct ggc gag 3237 Met Leu Ala Ser Gly Pro Ile Glu Val Gly Ser Leu Pro Ala Gly Glu 950 955 agg gtg tcc cga ccc gaa gct gct gtg cct ggg tac cag cat cca agt 3285 Arg Val Ser Arg Pro Glu Ala Ala Val Pro Gly Tyr Gln His Pro Ser 965 970 975 gaa gct atg aat gcc cac aca ccc cag ttc ttc atc tat aaa att gat 3333 Glu Ala Met Asn Ala His Thr Pro Gln Phe Phe Ile Tyr Lys Ile Asp 980 985 990 tca tcc aac cga gag cag cgg cta gag gac aaa gga gac acc cca ctg 3381 Ser Ser Asn Arg Glu Gln Arg Leu Glu Asp Lys Gly Asp Thr Pro Leu 1000 1005 gag ctg ggt gac gac tgt agc ctg gct ctc gtc tgg cgg aac aat gag 3429 Glu Leu Gly Asp Asp Cys Ser Leu Ala Leu Val Trp Arg Asn Asn Glu 1015 1020 cgc ttg cag gag ttt gtg ttg gta gcc tcc aag gag ctg gaa tgt gct 3477 Arg Leu Gln Glu Phe Val Leu Val Ala Ser Lys Glu Leu Glu Cys Ala 1035 1030 1040 gag gat cca ggc tct gcc ggt gag gct gcc cgg gcc ggc cac ttc acc 3525 Glu Asp Pro Gly Ser Ala Gly Glu Ala Ala Arg Ala Gly His Phe Thr 1045 1050 ctq qac caq tqc ctc aac ctc ttc aca cqq cct qaq qtq ctq qca ccc 3573 Leu Asp Gln Cys Leu Asn Leu Phe Thr Arg Pro Glu Val Leu Ala Pro 1060 1065 1070 1075 gag gag gcc tgg tac tgc cca cag tgc aaa cag cac cgt gag gcc tcc 3621 Glu Glu Ala Trp Tyr Cys Pro Gln Cys Lys Gln His Arg Glu Ala Ser 1080 1090 aag cag ctg ttg cta tgg cgc ctg cca aat gtt ctc atc gtg cag ctc 3669 Lys Gln Leu Leu Trp Arg Leu Pro Asn Val Leu Ile Val Gln Leu 1095 1100 1105 aag ege tte tee ttt egt agt ttt ate tgg egt gae aag ate aat gae 3717 Lys Arg Phe Ser Phe Arg Ser Phe Ile Trp Arg Asp Lys Ile Asn Asp 1110 1115 1120

ttg gtg gag ttc cct gtt agg aac ctg gac ctg agc aag ttc tgc att Leu Val Glu Phe Pro Val Arg Asn Leu Asp Leu Ser Lys Phe Cys Ile 1125 1130 1135	3765
ggt cag aaa gag gag cag ctg ccc agc tac gat cta tat gct gtc atc Gly Gln Lys Glu Glu Gln Leu Pro Ser Tyr Asp Leu Tyr Ala Val Ile 1140 1145 1150 1155	3813
aac cac tat gga ggc atg att ggt ggc cac tac act gcc tgt gca cgc Asn His Tyr Gly Gly Met Ile Gly Gly His Tyr Thr Ala Cys Ala Arg 1160 1165 1170	3861
ctg ccc aat gat cgt agc agt cag cgc agt gac gtg ggc tgg cgc ttg Leu Pro Asn Asp Arg Ser Ser Gln Arg Ser Asp Val Gly Trp Arg Leu 1175 1180 1185	3909
ttt gat gac agc aca gtg aca acg gta gac gag agc cag gtt gtg acg Phe Asp Asp Ser Thr Val Thr Thr Val Asp Glu Ser Gln Val Val Thr 1190 1195 1200	3957
cgt tat gcc tat gta ctc ttc tac cgc cgg cgg aac tct cct gtg gag Arg Tyr Ala Tyr Val Leu Phe Tyr Arg Arg Arg Asn Ser Pro Val Glu 1205 1210 1215	4005
agg ccc ccc agg gca ggt cac tct gag cac cac cca gac cta ggc cct Arg Pro Pro Arg Ala Gly His Ser Glu His His Pro Asp Leu Gly Pro 1220 1235	4053
gca gct gag gct gct gcc agc cag gct tcc cgg att tgg cag gag ctg Ala Ala Glu Ala Ala Ala Ser Gln Ala Ser Arg Ile Trp Gln Glu Leu 1240 1245 1250	4101
gag gct gag gag ccg gtg cct gag ggg tct ggg ccc ctg ggt ccc Glu Ala Glu Glu Pro Val Pro Glu Gly Ser Gly Pro Leu Gly Pro 1255 1260 1265	4149
tgg ggg ccc caa gac tgg gtg ggc ccc cta cca cgt ggc cct acc aca Trp Gly Pro Gln Asp Trp Val Gly Pro Leu Pro Arg Gly Pro Thr Thr 1270 1275 1280	4197
cca gat gag ggc tgc ctc cgg tac ttt gtc ctg ggc acc gtg gcg gct Pro Asp Glu Gly Cys Leu Arg Tyr Phe Val Leu Gly Thr Val Ala Ala 1285 1290 1295	4245
ttg gtg gcc ctc gtg ctc aac gtg ttc tat cct ctg gta tcc cag agt Leu Val Ala Leu Val Leu Asn Val Phe Tyr Pro Leu Val Ser Gln Ser 1300 1315	4293
cgc tgg aga tga gct cgcctgcagg cagctgctgt gagctggcct acctgcctgc Arg Trp Arg *	4348
cccaggccat gcctgccttt gttgtgggga acacctctgg gctttgggcc tcagcttatg	4408
catctggtgg gagaggttgt gcccctgcag gggcagagta tcctagggtg	4468

tgtatccatc	tggctgtctg	tccattcatc	ctgctgctct	gacccttggc	ctcaggcttg	4528
gccctgccca	agctacttcc	tgtacttaaa	agtgttaata	aaaccagact	attcaggccc	4588
aaaaaaaaa	aa					4600

		<2: <2:	10> 3 11> 2 12> I 13> I	2663	sapi	iens											
			21> () (2	2157))										
	cgga		00> 3		cgac	ga ti	tcgt	cgc	g gc	egget	gag	tcct	ctc	cag d	ccgc	gagagg	60
Dar bed Ja ton ton Holf Kad Huly Univ mid their reak met ind time	cgti	ttct	cca t	ccg	egget	c go	egege	ctcg	c tct	tgago	ccc	cgc	gecea	agg t	ggg	atg Met 1	117
		gaa Glu															165
		aga Arg															213
		tac Tyr 35															261
		gat Asp															309
		gat Asp															357
		gaa Glu															405
		ttg Leu															453
		gaa Glu 115															501

tgg Trp 130	Asp	tgt Cys	gaa Glu	ttt Phe	atc Ile 135	Arg	ctg Leu	cat His	ttt Phe	999 Gly 140	His	aac Asn	cgg Arg	aag Lys	aag Lys 145	549
cat His	ctt Leu	aac Asn	tac Tyr	aca Thr 150	Glu	ttc Phe	acg Thr	cag Gln	ttt Phe 155	Leu	cag Gln	gag Glu	ctg Leu	caa Gln 160	ttg Leu	597
gaa Glu	cat His	gca Ala	aga Arg 165	caa Gln	gcc Ala	ttt Phe	gca Ala	ctc Leu 170	aaa Lys	gac Asp	aaa Lys	agc Ser	aaa Lys 175	Ser	ggc Gly	645
atg Met	att Ile	tct Ser 180	ggt Gly	ctg Leu	gat Asp	ttc Phe	agt Ser 185	gac Asp	atc Ile	atg Met	gtt Val	acc Thr 190	att Ile	aga Arg	tct Ser	693
cac His	atg Met 195	ctt Leu	act Thr	cct Pro	ttt Phe	gtg Val 200	gag Glu	gag Glu	aac Asn	tta Leu	gtt Val 205	tca Ser	gca Ala	gct Ala	gga Gly	741
gga Gly 210	agt Ser	atc Ile	tca Ser	cac His	cag Gln 215	gtt Val	agc Ser	ttc Phe	tcc Ser	tac Tyr 220	ttc Phe	aat Asn	gca Ala	ttt Phe	aac Asn 225	789
tcg Ser	tta Leu	ctg Leu	aat Asn	aac Asn 230	atg Met	gag Glu	ctt Leu	gtt Val	cgt Arg 235	aag Lys	ata Ile	tat Tyr	agc Ser	act Thr 240	cta Leu	837
gct Ala	ggc Gly	aca Thr	agg Arg 245	aaa Lys	gat Asp	gtt Val	gaa Glu	gtc Val 250	aca Thr	aag Lys	gag Glu	gaa Glu	ttt Phe 255	gcc Ala	cag Gln	885
agt Ser	gcc Ala	ata Ile 260	cgc Arg	tat Tyr	gga Gly	caa Gln	gtc Val 265	aca Thr	cca Pro	cta Leu	gaa Glu	att Ile 270	gat Asp	att Ile	cta Leu	933
tat Tyr	cag Gln 275	ctt Leu	gca Ala	gac Asp	tta Leu	tat Tyr 280	aat Asn	gct Ala	tca Ser	gly ggg	cgc Arg 285	ttg Leu	act Thr	ttg Leu	gca Ala	981
gat Asp 290	att Ile	gag Glu	aga Arg	ata Ile	gcc Ala 295	cca Pro	ttg Leu	gct Ala	gag Glu	300 Gly aaa	gcc Ala	tta Leu	cct Pro	tac Tyr	aac Asn 305	1029
ctg Leu	gca Ala	gaa Glu	ctt Leu	cag Gln 310	aga Arg	cag Gln	cag Gln	tct Ser	cct Pro 315	ggg Gly	tta Leu	ggc Gly	agg Arg	cct Pro 320	atc Ile	1077
tgg Trp	ctc Leu	cag Gln	att Ile 325	gcc Ala	gag Glu	tct Ser	gct Ala	tac Tyr 330	aga Arg	ttc Phe	act Thr	ctg Leu	ggc Gly 335	tca Ser	gtt Val	1125
gct Ala	gga Gly	gct Ala 340	gtg Val	gga Gly	gcc Ala	act Thr	gca Ala 345	gtg Val	tat Tyr	cct Pro	ata Ile	gat Asp 350	ctg Leu	gtg Val	aag Lys	1173
acc	cga	atg	caa	aac	cag	cgt	ggc	tct	ggc	tct	gtt	gtt	ggg	gag	cta	1221



590 585 580 ggt gtt acc ttg gtc act tat gaa ctt ctc cag cgg tgg ttt tac att 1941 Gly Val Thr Leu Val Thr Tyr Glu Leu Leu Gln Arg Trp Phe Tyr Ile 595 gat ttt gga ggc ctc aaa ccc gct ggt tca gaa cca aca cct aag tca 1989 Asp Phe Gly Gly Leu Lys Pro Ala Gly Ser Glu Pro Thr Pro Lys Ser 620 610 cgc att gca gac ctt cct cct gcc aac cct gat cac atc ggt gga tac 2037 Arg Ile Ala Asp Leu Pro Pro Ala Asn Pro Asp His Ile Gly Gly Tyr 635 630 aga ctc gcc aca gcc acg ttt gca ggc atc gaa aac aaa ttt ggc ctt 2085 Arg Leu Ala Thr Ala Thr Phe Ala Gly Ile Glu Asn Lys Phe Gly Leu 655 645 tat ctc ccg aaa ttt aag tct cct agt gtt gct gtg gtt cag cca aag 2133 Tyr Leu Pro Lys Phe Lys Ser Pro Ser Val Ala Val Val Gln Pro Lys gca gca gtg gca gcc act cag tga tgagacaact gttgagtgtg gcaaaatggc 2187 Ala Ala Val Ala Ala Thr Gln 675 gccttgaaga aagagcctag gagagcagcc ctgtaatgta tccagtcagc tgcatggtac 2247 tgactgagct gaggagtcaa actcttcttt ctgtatgaca tatacatata cttgtttata 2307 aaataatcat ttgcccaggg aaaaaaccac aacgctgttt caagctttag tcttatgtgt 2367 tgaaatgttt ttgtaagcct tggcatgaat tagtgttcta gactctgctt tgcacagctt 2427 gcacttacag tgattgtaca tattgtacat ctttgtacag agacatcttg gcacctcatc 2487 ccaacaaatc acatttgtag aaatgtaatg cggttctgag tggcttgaaa tgtacagaat 2547 gttttgaaag tgttttatta agaatcacac aaaaataaat gtattaaaat taaattcatt 2607 ctcttattgg tgacttatgg aaataaagca tcaatattgg atgtaaaaaa aaaaaa 2663

```
<210> 4
```

<211> 6833

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (19)..(2364)

<220>

<221> misc_feature

<222> (1) . . . (6833)

<223> n = a,t,c or g

<400> 4 cgcgaggcaa canatgad	e atg ttg go Met Leu A	cc ttc ctg t la Phe Leu S 5	cct ggg atg ccg Ser Gly Met Pro	gtg acc 51 Val Thr 10
aga aac acc aag ta Arg Asn Thr Lys Ty 15	c ctc gac ctc c Leu Asp Le	c aag aat to u Lys Asn Se 20	ca caa gag atg o er Gln Glu Met 1 25	ctc cgc 99 Leu Arg
tac aaa gag gtc tg Tyr Lys Glu Val Cy 30	tac tac at Tyr Tyr Me 3	t Leu Phe Al	cc ctg gct gcc t la Leu Ala Ala ' 40	cac ggg 147 Tyr Gly
tgg ccc atg tac ct Trp Pro Met Tyr Le 45	g atg cgg aa 1 Met Arg Ly 50	g ccc gcc to s Pro Ala Cy	gc ggc ctc tgc ys Gly Leu Cys (55	caa ctg 195 Gln Leu
gct cgg tcc tgc tc Ala Arg Ser Cys Se 60	g tgt tgc ct r Cys Cys Le 65	u Cys Pro Al	cg agg ccg cgg la Arg Pro Arg 70	ttc gcc 243 Phe Ala 75
cct gga gtc acc at Pro Gly Val Thr Il 8	e Glu Glu As	c aac tgc tg p Asn Cys Cy 85	gt ggc tgt aat ys Gly Cys Asn	gcc att 291 Ala Ile 90
gcc atc cgg cgc ca Ala Ile Arg Arg Hi 95	c ttc ctg ga s Phe Leu As	c gag aac a p Glu Asn Mo 100	tg act gcg gtg et Thr Ala Val 105	gac atc 339 Asp Ile
gtc tat acc tcc tg Val Tyr Thr Ser Cy 110	c cat gat go s His Asp Al 11	a Val Tyr G	aa acg ccc ttc lu Thr Pro Phe 120	tac gtg 387 Tyr Val
gcg gtg gac cat ga Ala Val Asp His As 125	c aag aag aa p Lys Lys Ly 130	a gtg gtg a s Val Val I	tc agt atc cgg le Ser Ile Arg 135	ggg acc 435 Gly Thr
ctg tcc ccc aag ga Leu Ser Pro Lys As 140	t gcc ctg ac p Ala Leu Th 145	ır Asp Leu T	cg ggt gat gct hr Gly Asp Ala 50	gag cgc 483 Glu Arg 155
ctc ccc gtg gag gg Leu Pro Val Glu Gl 16	y His His G	gc acc tgg c Ly Thr Trp L 165	tg ggc cac aag eu Gly His Lys	ggt atg 531 Gly Met 170
gtc ctc tca gct ga Val Leu Ser Ala G 175	g tac atc aa u Tyr Ile Ly	ag aag aaa c ys Lys Lys L 180	etg gag cag gag Leu Glu Gln Glu 185	atg gtc 579 Met Val
ctg tcc cag gcc t Leu Ser Gln Ala Pl 190	ne Gly Arg A	ac ctg ggc c sp Leu Gly A 95	egc gga acc aaa Arg Gly Thr Lys 200	cac tac 627 His Tyr
ggc ctg att gtg g Gly Leu Ile Val V	ng ggc cac to al Gly His S	cc ctg ggc g er Leu Gly A	geg gge act gct Ala Gly Thr Ala	gcc atc 675 Ala Ile

205 215 210 ctc tcc ttc ctt ctg cgc cca cag tat ccg acc ctc aag tgc ttt gcc 723 Leu Ser Phe Leu Leu Arg Pro Gln Tyr Pro Thr Leu Lys Cys Phe Ala 225 230 771 tac tcc ccg cca ggg ggc ctg ctg agt gag gat gcg atg gag tat tcc Tyr Ser Pro Pro Gly Gly Leu Leu Ser Glu Asp Ala Met Glu Tyr Ser 240 245 819 aag gag tte gtg act gtt gtg gtt ctg ggc aaa gac cte gte eec agg Lys Glu Phe Val Thr Ala Val Val Leu Gly Lys Asp Leu Val Pro Arg 255 260 867 att ggc ctc tct cag ctg gaa ggc ttc cgc aga cag ctc ctg gat gtc Ile Gly Leu Ser Gln Leu Glu Gly Phe Arg Arg Gln Leu Leu Asp Val 270 275 ctg cag cga agc acc aag ccc aaa tgg cgg atc atc gtg ggg gcc acc 915 Leu Gln Arg Ser Thr Lys Pro Lys Trp Arg Ile Ile Val Gly Ala Thr 285 290 aaa tgc atc ccc aag tcg gag ctg cct gag gag gta gag gtg acc acc 963 Lys Cys Ile Pro Lys Ser Glu Leu Pro Glu Glu Val Glu Val Thr Thr 300 ctg gcc agc acg cgg ctc tgg acc cac ccc agc gac cta act ata gcc 1011 Leu Ala Ser Thr Arg Leu Trp Thr His Pro Ser Asp Leu Thr Ile Ala 320 ctc tca gcc agc act cca ctc tac ccg ccc ggc cgc atc atc cac gtg 1059 Leu Ser Ala Ser Thr Pro Leu Tyr Pro Pro Gly Arg Ile Ile His Val 335 340 gtc cac aac cac cct gca gag cag tgc tgc tgc tgt gag cag gag gag 1107 Val His Asn His Pro Ala Glu Gln Cys Cys Cys Glu Gln Glu Glu 350 355 ccc aca tac ttt gcc atc tgg ggc gac aac aag gcc ttc aat gag gtg 1155 Pro Thr Tyr Phe Ala Ile Trp Gly Asp Asn Lys Ala Phe Asn Glu Val 365 370 atc atc tcq cca qcc atq ctq cat qaq cac ctq ccc tat qtq qtc atq 1203 Ile Ile Ser Pro Ala Met Leu His Glu His Leu Pro Tyr Val Val Met 380 385 390 395 gag ggg ctc aac aag gtg ctg gag aac tac aac aag ggg aag acc gct 1251 Glu Gly Leu Asn Lys Val Leu Glu Asn Tyr Asn Lys Gly Lys Thr Ala 400 410 ctg ctc tct gca gcc aag gtc atg gtg agc cct acc gag gtg gac ctg 1299 Leu Leu Ser Ala Ala Lys Val Met Val Ser Pro Thr Glu Val Asp Leu 415 act cet gag etc ate tte cag cag cag eca etc ecc acg ggg eeg ecc 1347 Thr Pro Glu Leu Ile Phe Gln Gln Pro Leu Pro Thr Gly Pro Pro 430 435

				gcc Ala							1395
_	_	_	 -	aag Lys 465	_		_	_	_	 	1443
				ctg Leu							1491
				ctg Leu							1539
				cgg Arg							1587
				tac Tyr							1635
				tcc Ser 545							1683
				tac Tyr							1731
				act Thr							1779
				cag Gln							1827
				ccc Pro							1875
				ggt Gly 625							1923
				ctg Leu							1971
				gac Asp							2019

														acc Thr		2067
														ctg Leu		2115
														ctc Leu		2163
														tcc Ser 730		2211
														ggc Gly		2259
agt Ser	agc Ser	cag Gln 750	gaa Glu	tgc Cys	ctg Leu	gcg Ala	gct Ala 755	gac Asp	aag Lys	atc Ile	cgg Arg	act Thr 760	tct Ser	acc Thr	ccc Pro	2307
														atc Ile		2355
gca Ala 780		tag *	caco	cca	gttg	gegte	igc c	cagco	gggd	cc ca	aggca	aggag	g cag	ggtgg	lccc	2411
tgtg	ggca	acc t	ggtg	gccts	je ec	ccts	ccgg	g gca	ıgctt	taa	ggad	agac	cc d	ccago	ggcag	2471
ttta	ıgcct	ca g	ggcac	aggo	a to	gctg	ıctga	gct	gggg	gtc	cgca	tccc	cta d	cctca	ıgctta	2531
ggad	cccc	ag a	agcca	aggt	g go	tggg	gatct	ggc	ccca	ıcag	atgg	ggaa	ag a	tggg	gaagg	2591
gtgt	ggag	gtg g	gggag	gago	c to	ıggca	gcct	gct	gggt	:999	ccac	acto	cag o	ctga	ctgcc	2651
ctcc	atgg	199 S	gcatt	ctgg	jc ac	cccc	tgct	cca	ıggac	agg	ccat	gggc	aa g	gctgc	ctccc	2711
atca	ctgo	ct ç	gctgg	gctgc	t ct	ссса	9999	g cca	ıggtg	gag	agca	ıgtgo	cc c	ccga	cacat	2771
gtat	tctc	at c	ctgtg	gtcc	a gg	ccgg	cato	gto	ctgg	cca	cccc	ccag	gat o	tggt	gcctg	2831
ctgg	ccgg	jcc c	cctg	ıgggt	g co	cctg	ccga	ggt	ggcc	tgc	agtg	ctgt	ac a	tgtt	tacag	2891
aago	tgct	gg g	gcttg	gcto	a gg	atgt	gttc	tgg	gctt	gca	agco	cccc	gc c	caat	catgt	2951
gtto	agta	igc c	catco	tctg	a go	aggg	ссса	agg	cago	cag	gggd	ctgg	ag c	ggcc	agagg	3011
aggg	tggg	ıgt c	aggg	ccgc	c cc	ttct	ctgc	ctt	gtgc	ctc	tcat	gctg	rcc t	cctc	tgccc	3071
atgg	gtcc	tg g	gcac	ccag	ig co	tgcc	ctgo	ctg	ctgg	cta	cttc	ctgg	ict t	acct	tctac	3131
cccc	aagg	at c	ctca	ccac	сса	aagg	gtgg	tgg	gcac	tgc	tgtg	acca	cc c	cago	tgcag	3191

agtcagtgcc ctgggtggaa ggaaggcact gagagccccc ttcctctgag ggccccacct 3251 caccccttgg tgtcaccccc accacgccta ggcagctctg ggccctggga tctggaacca 3311 acacaccet gtteccetca getttecete etegetggee tgggeaccet eetgggagea 3371 ggccttcctc cctcccaccc ccaatgtcct gttggtagga ggtggggcca agagtggggt 3431 atggtgggcc ttggctggag acctctgtcc actgcccagg gaggggcctg gggctgggag 3491 cagtcccggt ttagcctgag gtccccatag ggcttcctcc cctgctgggt ttgggaagca 3551 gttagggaga tagcgacccg gagtttcccc agaagcgggg tgggagggtg tgcatgctag 3611 tgttggcgcg tatgcatgtg catgagtgtg caccgttcct aaggaagggg cctctggggc 3671 tgcccaccct acctgccctg cctgcctgct gcccctccca gcctgccaag aaaacggtag 3731 gggagcatga tggggccttt gaggcagggt cgcagggaca agctcagctt taggcaccat 3791 ctgttcccat cgcgcctgct gctgtgaccc gttttggaaa actggtgtgt accgaggcgc 3851 tgactgcacg gctgaccgcc tgctcgtgcc ttcattctgc agcggcatgg tccctcccat 3911 totggctcca cotgcagoot cootgggtgg cotaggotco cocgaccaag agacotcoot 3971 ctcatgatca ctggtacctg ggggcctgaa ttctggcccc cggctcccca cacagctggg 4031 actggcctgg atggctgtcc tgggagcccc tgcccaccct gacagaggga gctgggcctc 4091 ccctcatcct ctgtaactcc cgccttcacc agactcaagg acaccctggc cctgctgagg 4151 catacagage tteageeeag cacagaagea agacaaaate agtggetett agagtttaga 4211 aaacaagaca gactctcaga tgaaagatct gacaagcacc gtggccagtc acagggagag 4271 acttgatgtc tggcctttta attcctcctc tgccagggtg ggtcctggga cctctaatgt 4331 gggcatgtcg tccaccccag gacaagccat cagggacaga cccccaccc ccaaggctgc 4391 agccacacca tgtttcaggc ttggggctgg ggcaggcttg ggctcaatcc tgggcaccca 4451 ggggcagccc accectaacc tggctcctac ccaccttgcc cttgaaggat gggcctgctg 4511 cacgtctccc tcctccaccc cataccacac tggggggtct gagccacccc cctcagcccc 4571 gttcggctca gaccgacccc cactccatcc ccagacctgc agcacaagtg cgcgggcctg 4631 tecteccagg ggeetgggeg actecatatg caatcagtag egageageeg ggeeccacag 4691 acceteatge actetettae gtgccattet ecceagaett tttttgtaet taatgtatga 4751 aagatccaaa ctaatattgc tgtaaaaaagg agagacaaat taatatagct tattctataa 4811 atatatetgt atataaaggt ttetgtatat tgtatagage tgtgtataaa etggatgtag 4871

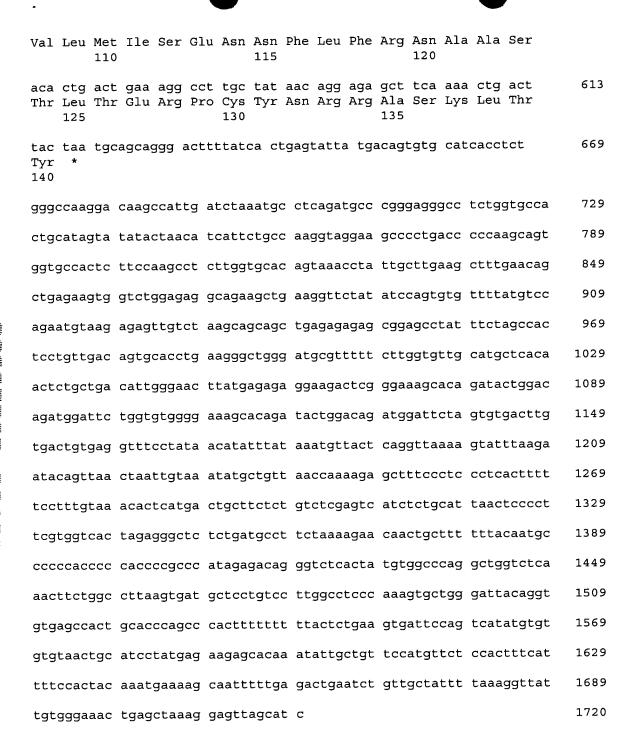
4931 aagcacgctg gctgcctcga atgtctttgc atcaggtggg gactgggtag aaatttgatg 4991 tegaggttge ageagageag ggggttggea tggggeeggg ggeegggggg geetetgeea ctgttccctg gatggaacag aaagcctgct cctgctcggc tagtgccctg gcccggggcc 5051 acactttgag ttgcagggag ggagtagaag acccttgaag cccctgtga atgagggcag 5111 5171 ccacttgcgg agtcctgccc cactttgagc cctcctcttc ctctgcgaaa tggcctgatg 5231 ccagtgctgt gtgggtccac aggaggccag gagagtccca cggtggggag gacagggctg 5291 taccttctcc tgggctggcc tctacgccca ttacccatta accctcaggt gccagcatcc 5351 ctctccccag tgctgccttc ggtctaccac ctcctcctgg ccctgctccc actcaaggga 5411 cagtgatggg tgctgagagc tgattggaac tggagagggc accatttact gatcactgac 5471 ctggcacttt acctccactg taaggcaggg atactgagtg tgctttatag atgaggcctc tgaggacaga gggcaggcct tggggactag gtggagctgg ctacagggga cagccatgtc 5531 tgctgggcta gggctgaaat ctcagcccct cactcactgt ggctcggtag agaagccagg 5591 ggcacagatg aggactcatc tccattgatg ggccccccta ggtccttgta tgcaagtccc 5651 ctgggctact ttcaccggcc cagccacctt cctgccccag cctctgcccc agcgctgctt 5711 5771 gggacccaac ttcattatgg agttggcaga tggcagcctc aactcttggc tgagcccttg agtotgggac atttcagcca cototttoot coagtocaga gatgaaaato cotggggaca 5831 gttgctcctt tgctcagtga cctagtgtaa caggggagat ggcagggcct gagctccctt 5891 5951 agccaggtca gctgctacag ggttaacagg aggctccatt ccaccccttc caacttcaag gctaccctag agattgaata atctatactc ttaattgatt ataatgcaat ggagttgggg 6011 6071 cgttagggac aaagtacgag tcttcccttc tacctccagc ccttgctgac caggacaggg 6131 acaatgtgta gctcaacgga tggtggtgag aacttagatg atggtcagtt atgcagtatg tgggatacgg aggaaagatc cgtgggtatg tggaggcttg tagagaagct ggttctgtgg 6191 6251 ctggtcccag gcgactcgta atgtaaatcc gtttctcaga atcgcgtggt gtaggcgggt 6311 gtctactttg tccgcaggca ggcctgaccc cgggtggagg aggggcaggg tggagaataa caattgtett aagggagtet geaagaeagg agggggtgge agagaagaac eteagetetg 6371 aagaagetea etgeeeagee etteeeacet teetetteae ggaeetagea eetteetggg 6431 6491 cctcagtttc tctcattgcc ttgggtgctg gagtatgtgg ggcctcctct cctatctcca ggccttcagc cccggctgcc acggggtgtg ggtacctctt ggttgggtct cggggtagga 6551 6611 tgatgtaatg gttctgtgca ttcgccagcg agggcagctg gggtctgttc ctagctctcc

tgettaccea cagtgettet ettggeegta teagggeace getgtgeete egetttetea 6671
tettgaagat taetggteee cagggtaggt cagtgeecet aagettaggg ggettgttga 6731
geatgttetg tggttetgtg tgeaaggeet gaaceatgae agetetggee cagegtggee 6791
tggteetggt eeetggeaca teeagtgggg eeegeeecae et 6833

<210> 5 <211> 1720 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (197)..(619) <400> 5 atgategeet teggaacegg ceeggaatte cegggtegae eeaegegtee getetttgge actagttcag aatggtgatg tgtcggcccc ctctgccata ctcagaacac cagaaagcac 120 aaaaccgggt cctgtttgtc agccaccagt gagtcagagc cgctccctgt tttcttctgt 180 229 cccqtccaaq ccacca atg tct ctg gag cct caa aat ggg acg tat gca Met Ser Leu Glu Pro Gln Asn Gly Thr Tyr Ala gga cca gcg cca gca ttc cag cca ttt ttc ttc act gga gca ttt cca 277 Gly Pro Ala Pro Ala Phe Gln Pro Phe Phe Thr Gly Ala Phe Pro ttt aat atg caa gag ctg gta ctc aag gtg aga att cag aac cca tct 325 Phe Asn Met Gln Glu Leu Val Leu Lys Val Arg Ile Gln Asn Pro Ser ctt cga gaa aat gat ttc att gaa att gaa ctg gac cga cag gag ctc 373 Leu Arg Glu Asn Asp Phe Ile Glu Ile Glu Leu Asp Arg Gln Glu Leu 45 50 acc tac caa gag ttg ctc aaa gtg tgt tgc tgt gag ctg ggt gtt aat 421 Thr Tyr Gln Glu Leu Leu Lys Val Cys Cys Glu Leu Gly Val Asn 60 75 cca gat caa gtg gag aag atc aga aag tta ccc aat act ctg tta agg 469 Pro Asp Gln Val Glu Lys Ile Arg Lys Leu Pro Asn Thr Leu Leu Arg 80 85 aag gac aag gat gtt gct cga ctc caa gat ttc cag gag ctg gaa ctg 517 Lys Asp Lys Asp Val Ala Arg Leu Gln Asp Phe Gln Glu Leu Glu Leu 95 100

565

gtt ctg atg ata agt gaa aat ttt ctg ttc aga aat gct gca tcc



<210> 6

<211> 1658

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (523)..(1479)

<400> 6 ctagtacgcc tgtggaacgc ctgcaggtac cggtccggaa ttcccgggtc gacccacgcg	60
tccgcccacg cgtccgctct gtctcaaata ataataataa taataataat aataataata	120
ataataataa tgtaggggac ttgatgaagg gaaaggatca gatagattct gaaaagaagt	180
acagaacctg ctaaggccat catacctatt gatcggaagt cagtccatca gatttgctct	240
	300
ggaccggtgg tactgagtct aatcactgcg tgtaagaaga tagtaggaaa cattctggat	
gctaggtgcc agtaatattg atctaaagct taaggactat ggaatggatc tcattgaagt	360
ttcaggcaat ggatgtgggg tagaagaaga aaacttcgaa ggcttaatct ctttcagctc	420
tgaaacatca cacatctaag attcaagagt ttgccgacct aactcgggtt gaaacttttg	480
gctttcgggg gaaagctctg agctcacttt gtgcactgag tg atg tca cca ttt Met Ser Pro Phe 1	534
cta cct gcc acg tat cgg cga agg ttg gga ctc gac tgg tgt ttg atc Leu Pro Ala Thr Tyr Arg Arg Leu Gly Leu Asp Trp Cys Leu Ile 5 10 15 20	582
acg atg gga aaa tca tcc aga aaa ccc cct acc ccc acc cca gag gga Thr Met Gly Lys Ser Ser Arg Lys Pro Pro Thr Pro Thr Pro Glu Gly 25 30 35	630
ccc aca gtc agc gtg aag cag tta ttt tct acg cta cct gtg cgc cat Pro Thr Val Ser Val Lys Gln Leu Phe Ser Thr Leu Pro Val Arg His 40 45 50	678
aag gaa ttt caa agg aat att aag aag aaa cgt gcc tgc ttc ccc ttc Lys Glu Phe Gln Arg Asn Ile Lys Lys Lys Arg Ala Cys Phe Pro Phe 55 60 65	726
gcc ttc tgc cgt gat tgt cag ttt ctt gag ggc tcc cca gcc atg ctt Ala Phe Cys Arg Asp Cys Gln Phe Leu Glu Gly Ser Pro Ala Met Leu 70 75 80	774
cct gta cag cct gca aaa ctt åca gaa cct gct aag gcc atc aaa cct Pro Val Gln Pro Ala Lys Leu Thr Glu Pro Ala Lys Ala Ile Lys Pro 85 90 95 100	822
att gat cgg aag tca gtc cat cag att tgc tct ggg ccg gtg gta ctg Ile Asp Arg Lys Ser Val His Gln Ile Cys Ser Gly Pro Val Val Leu 105 110 115	870
agt cta agc act gcg gtg aag aag ata gta gga aac agt ctg gat gct Ser Leu Ser Thr Ala Val Lys Lys Ile Val Gly Asn Ser Leu Asp Ala 120 125 130	918
ggt gcc act aat att gat cta aag ctt aag gac tat gga atg gat ctc Gly Ala Thr Asn Ile Asp Leu Lys Leu Lys Asp Tyr Gly Met Asp Leu	966

		135					140					145				
att Ile	gaa Glu 150	gtt Val	tca Ser	ggc Gly	aat Asn	gga Gly 155	tgt Cys	gly ggg	gta Val	gaa Glu	gaa Glu 160	gaa Glu	aac Asn	ttc Phe	gaa Glu	1014
											tct Ser					1062
											ttt Phe					1110
											att Ile					1158
											gat Asp					1206
											999 Gly 240					1254
											cat His					1302
											ttc Phe					1350
											ctt Leu					1398
gca Ala	aaa Lys	ctg Leu 295	act Thr	gta Val	act Thr	gga Gly	gag Glu 300	cta Leu	cgg Arg	gca Ala	tgc Cys	aga Arg 305	agt Ser	tgg Trp	aag Lys	1446
acg Thr	agg Arg 310	gaa Glu	ggc Gly	atc Ile	aca Thr	gag Glu 315	gct Ala	gtg Val	999 Gly	tga *	accg	gactt	caa	ıggaa	ıtgg	1497
gtcc	ttcc	ct t	caga	acca	ıc at	gtgt	gcgg	gac	acco	aga	caga	aaac	ac a	aatg	caaag	1557
tcaa	gtgg	gag g	gcat	ttgg	ja ag	gago	agtg	aag	ıccaa	.gcc	agga	aaca	ıcc a	agat	ggcga	1617
tcaagtggag ggcatttgga aggagcagtg aagccaagcc											1658					

<210> 7 <211> 3739 <212> DNA <213> Homo sapiens

<220> <221> CDS

<222> (460) .. (3621)

<400> 7 gtctgctgta ataccctcta ctatagggac cactttgtac aagaaagctg ggtacgcgta 60 120 agettgggcc cetegaggga ttetetagag egteegeggg ggetgeeagg gtattteggg aagggggcgt gaggaggcgg cggcggcagc ggcgggtagg gcaggcagca gagggaagga 180 240 gaaagaaagg aaggaagag gcggggagtc ctcagaggag gaggcgggac cggccgggca 300 geogeceet geoegegte tteteagege agtegggege ggaceegetg gteeegggea 360 gcggccaagg ctactggggc gggagcagtg ggccggtcgg cggcggcagc ggcagcggcg 420 474 atg gcg ttc ctc aaa gaggaggagg aggctggagt gggcgcggag gcgaccgcc Met Ala Phe Leu Lys 5 522 ctc cgt gac cag cca tca ctg gtg caa gct ata ttt aac gga gat cct Leu Arg Asp Gln Pro Ser Leu Val Gln Ala Ile Phe Asn Gly Asp Pro 570 gat gaa gtt cga gca cta ata ttt aag aaa gaa gat gtt aac ttt cag Asp Glu Val Arg Ala Leu Ile Phe Lys Lys Glu Asp Val Asn Phe Gln 618 gac aat gaa aag cga acc cca ttg cac gcc gca gct tac ctt gga gat Asp Asn Glu Lys Arg Thr Pro Leu His Ala Ala Ala Tyr Leu Gly Asp 45 40 gca gaa atc att gaa ctt ctt att tta tct gga gct aga gtt aat gcc 666 Ala Glu Ile Ile Glu Leu Leu Ile Leu Ser Gly Ala Arg Val Asn Ala 55 60 aaa gac agc aaa tgg ttg aca cct tta cac aga gca gtt gca tct tgt 714 Lys Asp Ser Lys Trp Leu Thr Pro Leu His Arg Ala Val Ala Ser Cys 75 85 70 agt gag gaa gca gtt cag gta ctt ttg aag cat tct gca gat gtt aat 762 Ser Glu Glu Ala Val Gln Val Leu Leu Lys His Ser Ala Asp Val Asn 100 90 gct cga gac aaa aat tgg caa acc cct tta cat ata gct gct gct aat 810 Ala Arg Asp Lys Asn Trp Gln Thr Pro Leu His Ile Ala Ala Asn 110 115 105 aaa gct gta aag tgt gct gaa gct ttg gta cct ctt ctg agt aat gta 858 Lys Ala Val Lys Cys Ala Glu Ala Leu Val Pro Leu Leu Ser Asn Val

125

aac Asn	gta Val 135	tct Ser	gat Asp	cga Arg	gca Ala	999 Gly 140	agg Arg	act Thr	gca Ala	tta Leu	cat His 145	cat His	gca Ala	gct Ala	ttc Phe	906
agt Ser 150	gga Gly	cat His	ggt Gly	gag Glu	atg Met 155	gtc Val	aaa Lys	cta Leu	ctc Leu	ttg Leu 160	tct Ser	aga Arg	ggt Gly	gcc Ala	aat Asn 165	954
att Ile	aat Asn	gct Ala	ttt Phe	gac Asp 170	aag Lys	aaa Lys	gat Asp	agg Arg	cgt Arg 175	gct Ala	atc Ile	cat His	tgg Trp	gca Ala 180	gca Ala	1002
tat Tyr	atg Met	ggt Gly	cac His 185	att Ile	gaa Glu	gta Val	gtg Val	aaa Lys 190	ttg Leu	ctt Leu	gtg Val	tcg Ser	cat His 195	gga Gly	gct Ala	1050
gaa Glu	gtg Val	aca Thr 200	tgc Cys	aag Lys	gat Asp	aaa Lys	aag Lys 205	tct Ser	tat Tyr	aca Thr	cct Pro	ctt Leu 210	cat His	gca Ala	gca Ala	1098
gcc Ala	tct Ser 215	agt Ser	gga Gly	atg Met	atc Ile	agc Ser 220	gta Val	gtc Val	aag Lys	tac Tyr	ctt Leu 225	cta Leu	gat Asp	ctt Leu	gga Gly	1146
gtt Val 230	gat Asp	atg Met	aat Asn	gaa Glu	cca Pro 235	aat Asn	gcc Ala	tat Tyr	gga Gly	aat Asn 240	aca Thr	cct Pro	ctt Leu	cat His	gta Val 245	1194
gcc Ala	tgc Cys	tat Tyr	aat Asn	gga Gly 250	caa Gln	gat Asp	gtt Val	gta Val	gtg Val 255	aat Asn	gaa Glu	ctt Leu	ata Ile	gac Asp 260	tgt Cys	1242
ggt Gly	gct Ala	att Ile	gtg Val 265	aat Asn	caa Gln	aag Lys	aat Asn	gaa Glu 270	aaa Lys	gga Gly	ttt Phe	act Thr	cct Pro 275	ttg Leu	cac His	1290
ttt Phe	gct Ala	gct Ala 280	gca Ala	tca Ser	aca Thr	cat His	gga Gly 285	gca Ala	ttg Leu	tgt Cys	tta Leu	gag Glu 290	ctt Leu	cta Leu	gtt Val	1338
ggc Gly	aat Asn 295	ggg Gly	gcc Ala	gat Asp	gtc Val	aat Asn 300	atg Met	aag Lys	agt Ser	aaa Lys	gat Asp 305	Gly	aaa Lys	acc Thr	cca Pro	1386
cta Leu 310	cac His	atg Met	act Thr	gct Ala	ctc Leu 315	cac His	ggt Gly	aga Arg	ttc Phe	tcc Ser 320	cga Arg	tca Ser	caa Gln	acc Thr	att Ile 325	1434
atc Ile	cag Gln	agt Ser	gga Gly	gct Ala 330	gta Val	atc Ile	gac Asp	tgt Cys	gag Glu 335	gat Asp	aag Lys	aat Asn	gga Gly	aat Asn 340	acc Thr	1482
cct Pro	ttg Leu	cac His	ata Ile 345	Ala	gca Ala	cgg Arg	tat Tyr	ggc Gly 350	His	gag Glu	ctg Leu	ctg Leu	atc Ile 355	Asn	act Thr	1530

ctt Leu	att Ile	aca Thr 360	agt Ser	ggt Gly	gct Ala	gac Asp	act Thr 365	gca Ala	aag Lys	cgt Arg	ggc	ata Ile 370	cat His	gga Gly	atg Met	1578
ttc Phe	ccc Pro 375	ctc Leu	cat His	ttg Leu	gca Ala	gcc Ala 380	tta Leu	agc Ser	ggc Gly	ttt Phe	tca Ser 385	gat Asp	tgc Cys	tgc Cys	aga Arg	1626
aaa Lys 390	ctt Leu	ctt Leu	tct Ser	tca Ser	gga Gly 395	ttt Phe	gat Asp	ata Ile	gat Asp	acc Thr 400	cca Pro	gat Asp	gat Asp	ttt Phe	ggc Gly 405	1674
agg Arg	act Thr	tgt Cys	cta Leu	cat His 410	gca Ala	gct Ala	gca Ala	gct Ala	gga Gly 415	ggg Gly	aat Asn	ttg Leu	gag Glu	tgc Cys 420	cta Leu	1722
aac Asn	ctt Leu	ctg Leu	ctg Leu 425	aat Asn	act Thr	ggt Gly	gca Ala	gac Asp 430	ttt Phe	aat Asn	aaa Lys	aag Lys	gac Asp 435	aaa Lys	ttt Phe	1770
Gly 999	aga Arg	tct Ser 440	cca Pro	ctg Leu	cac His	tac Tyr	gct Ala 445	gct Ala	gcc Ala	aac Asn	tgc Cys	aat Asn 450	tac Tyr	cag Gln	tgc Cys	1818
ctg Leu	ttt Phe 455	gct Ala	ctt Leu	gtg Val	gga Gly	tca Ser 460	gga Gly	gca Ala	agt Ser	gtg Val	aat Asn 465	gac Asp	ctt Leu	gat Asp	gaa Glu	1866
aga Arg 470	ggc Gly	tgc Cys	aca Thr	ccc Pro	ctg Leu 475	cac His	tat Tyr	gca Ala	gct Ala	aca Thr 480	tca Ser	gac Asp	aca Thr	gat Asp	ggc Gly 485	1914
Lys	Cys	Leu	Glu	Tyr 490		Leu	Arg	Asn	495	Ala	Asn	Pro	Gly	11e 500	Arg	1962
gat Asp	aag Lys	caa Gln	gga Gly 505	tac Tyr	aac Asn	gca Ala	gtt Val	cat His 510	Tyr	tca Ser	gct Ala	gct Ala	tat Tyr 515	GIY	cac	2010
cgt Arg	cta Leu	tgt Cys 520	Leu	cag Gln	ctg Leu	att Ile	gca Ala 525	Ser	gaa Glu	act Thr	cct	cta Leu 530	Asp	gtt Val	tta Leu	2058
atg Met	gaa Glu 535	Thr	tca Ser	gga Gly	aca Thr	gac Asp 540	Met	ctg Leu	g agt Ser	gat Asp	tca Ser 545	Asp	aat Asn	aga Arg	gca Ala	2106
aca Thr 550	Ile	ago Ser	cct Pro	tta Leu	cac His 555	Leu	gct Ala	gcc Ala	tat Tyr	cat His 560	Gly	cac His	cat His	caa Glr	gca Ala 565	2154
ctg Leu	gaa Glu	ı gtç ı Val	ttg Lev	gta Val 570	. Gln	tct Ser	ttg Lev	g tta 1 Leu	a gat a Asp 575	Leu	gat Asp	gto Val	aga Arç	aat JAsr 580	agt Ser	2202
agt	gga	aga	a aca	ccc	c cta	gat	ctt	gca	a gct	ttt	aag	g ggd	cat	gtt	gaa	2250

Ser Gl	Ly Ai		hr 1 85	Pro	Leu	Asp	Leu	Al 59	.a A 90	la 1	Phe	Lys	; Gl	Ly E	lis 595	Val	G1	.u	
tgt gt Cys Va	al A	at g sp V 00	ta al	ctc Leu	att Ile	aat Asn	cag Glr 605	1 63	ga g Ly A	cc la	tca Ser	ato Ile		ta q eu ' 10	gta Val	aaa Lys	ga As	at sp	2298
tac at Tyr I	tt t le L 15	tg a eu I	lag Lys	agg Arg	aca Thr	cct Pro 620	TTC	c ca	at g is A	ca la	gca Ala	gca Ala 62!		ca hr	aat Asn	ggt Gly	Ca H:	at is	2346
tca g Ser G 630		gc t	tta Leu	cgg Arg	cta Leu 635	tta Leu	ata Il	a g e G	ga a ly <i>P</i>	at Asn	gca Ala 640	ga Gl	a c u P	ca ro	cag Gln	aat Asn	9 A 6	ca 1a 45	2394
gtg g Val A	jat a Asp]	att (caa Gln	gat Asp 650	gga Gly	aat Asr	gg Gl	a c y G	TIL	acg Thr 655	cct Pro	ct Le	g a u M	itg Iet	cta Leu	tct Ser 660	; g ; V	tt al	2442
ctc a Leu A	aac (Asn (Gly	cac His 665	aca Thr	gac Asp	tgt Cy	t gt s Va	т т	ac Yr 570	tca Ser	ttg Leu	g ct Le	ga eu P	aac Asn	aaa Lys 675	gga	a g	jca Ala	2490
aat (Asn)	Val .	gat Asp 680	gcc Ala	aaa Lys	gat Asp	aa Ly	g to s Ti	.р С	gga Bly	agg Arg	aca Thr	a go c Al	eg 1 la 1	ttg Leu 690	cat His	age Are	a g g (31y 999	2538
gca (ggc	cat His	ga:	a ga u Gl 70	u C	gt (ys '	gta Val	gat Asp	gca Ala		ta eu 05	ctt Leu	caa Glr	a ca n Hi	t (ggt Gly	2586
gct Ala 710		tgc Cys	tta Leu	a ct	t cg u Ar 71	g As	ıt a sp S	gc er	agg Arg	ggc Gly	cg Ar	9 -	cg hr	cct Pro	ata Ile	a ca e Hi	.s	ctg Leu 725	2634
tct Ser	gct Ala	gcc Ala	tgt Cy:	= gg = Gl 73	у ні	c at	t g le G	gt ly	gtt Val	ctt Let 73!		a g y A	cc la	ctt Lei	tte	g ca u Gl 74	ag Ln 10	tca Ser	2682
gca Ala	gca Ala	tct Ser	ate Me	t As	it go	a aa la A	at c sn F	ca Pro	gcc Ala 750	111	a go r Al	a g La <i>P</i>	ac Asp	aat Ası	ca n Hi 75	t gg s G:	ga ly	tat Tyr	2730
acg Thr	gca Ala	ctt Leu 760	ı Hi	c to	rp A	ct t la C	ys .	ac Tyr 765	aat Asn	gg Gl	t ca y Hi	ac g is (gag Glu	ac Th 77	a tg r Cy 0	jt g vs V	ta al	gaa Glu	2778
ctg Leu	ctt Leu 775	Let	a ga ı Gl	a ca .u G	ag g ln G	⊥u ∨	tt i	ttc Phe	caç Glr	g aa n Ly	a ao 's Tì	***	gaa Glu 785		a aa y As	at g sn A	ct la	ttt Phe	2826
agt Ser 790	cca Pro		g ca u Hi	at t is C	ys A	cc s la V 95	gtg /al	ata Ile	aat Asi	t ga n As	,p	ac sn	gaa Glu	ı gg	jt go .y A:	ct g la A	jct Ala	gag Glu 805	2874
		a at u Il	t ga e A	at a sp T	ca t	ta q eu (ggt Gly	gcc Ala	ag Se	c at	t g le V	ıtg Val	aac Asr	c go	cc a la T	ca g hr <i>l</i>	gat Asp	tca Ser	2922

aaa gga aga act cct ctc cat gca gcc gcc ttc aca gac cat gta gag Lys Gly Arg Thr Pro Leu His Ala Ala Phe Thr Asp His Val Glu tgt tta cag ctg ctc agc cat aat gct caa gtc aat tct gtg gac Cys Leu Gln Leu Leu Ser His Asn Ala Gln Val Asn Ser Val Asp tct aca ggg aaa aca cct ctt atg atg gct gca gaa aat gga caa aca Ser Thr Gly Lys Thr Pro Leu Met Met Ala Ala Glu Asn Gly Gln Thr aat aca gtt gag atg ctg gtt agc agt gct agt gca gaa ctg act tta Asn Thr Val Glu Met Leu Val Ser Ser Ala Ser Ala Glu Leu Thr Leu caa gat aac agt aaa aat act gcc ctc cat ttg gct tgt agc aag ggt Gln Asp Asn Ser Lys Asn Thr Ala Leu His Leu Ala Cys Ser Lys Gly cat gaa act agt gcc ttg tta ata ctg gaa aag ata aca gat aga aac His Glu Thr Ser Ala Leu Leu Ile Leu Glu Lys Ile Thr Asp Arg Asn ctc atc aat gca acc aac gca gcc ttg caa aca cct ctg cat gtt gct Leu Ile Asn Ala Thr Asn Ala Ala Leu Gln Thr Pro Leu His Val Ala gcc cga aat ggg cta aca atg gtg gtt cag gaa ctt ttg gga aaa gga Ala Arg Asn Gly Leu Thr Met Val Val Gln Glu Leu Leu Gly Lys Gly gca agt gtg ctt gca gta gat gaa aat ggc tat acc cca gct ttg gcc Ala Ser Val Leu Ala Val Asp Glu Asn Gly Tyr Thr Pro Ala Leu Ala tgt gct ccc aat aag gat gtg gct gat tgc ctg gct ctc att ttg gcc Cys Ala Pro Asn Lys Asp Val Ala Asp Cys Leu Ala Leu Ile Leu Ala acc atq atq cct qtc tca tca aqt aqt cct tta tca tcc tta aca ttc Thr Met Met Pro Val Ser Ser Ser Pro Leu Ser Ser Leu Thr Phe aat gcc att aac cgt tat acc aac acc tca aaa aca gtc agc ttt gaa Asn Ala Ile Asn Arg Tyr Thr Asn Thr Ser Lys Thr Val Ser Phe Glu gct ttg ccc atc atg agg aat gaa cct agc tcc tat tgc agt ttc aat Ala Leu Pro Ile Met Arg Asn Glu Pro Ser Ser Tyr Cys Ser Phe Asn aac att gga ggg gaa cag gag tac tta tac act gac gtg gat gag ctc Asn Ile Gly Gly Glu Gln Glu Tyr Leu Tyr Thr Asp Val Asp Glu Leu

aac gac tcc gat tct gag acc tac tga gaggc tgaggaggag ggagttctca Asn Asp Ser Asp Ser Glu Thr Tyr * 1050	3646
cagtaaagct tcaaactgtg ctttttcagg aaacaggcac tttgatattc acgtagaaat	3706
tcaacctaag agggacagtt ccgttgaagc cgt	3739
<210> 8 <211> 2162 <212> DNA <213> Homo sapiens	
<220> <221> CDS <222> (387)(1748)	
<400> 8 cgtactgaag tggtctgcat gagcgaccgg tccggaattc ccgggtcgac gatttcgttc	60
cggttgcact cttcctatag cccagagggc gagagggcct gtggcctggg ggaaggagga	120
cgaggttctg cctggatccc agcagtagga cgctgtgcca tttgggaaca aaggaatagt	180
ctgcctggaa tccctgcaga tcttggggcc ggaggccagt ccaacccttg gagcaggaag	240
aaacgcaaag ttgtcaagaa ccaagtcgag ctgcctcaga gccggcccgc agtagctgca	300
gacteegeee gegaegtgtg egegettete tgggeeagag egageetgtt ttgtgetegg	360
gttaagagat ttgtcccagc tatacc atg ggc cgc act cgg gaa gct ggc tgc Met Gly Arg Thr Arg Glu Ala Gly Cys 1 5	413
gtg gcc gct ggt gtg gtt atc ggg gct ggt gcc tgc tac tgt gta tac Val Ala Ala Gly Val Val Ile Gly Ala Gly Ala Cys Tyr Cys Val Tyr 10 15 20 25	461
aga ctg gct tgg gga aga gac gag aac gag aaa atc tgg gac gaa gac Arg Leu Ala Trp Gly Arg Asp Glu Asn Glu Lys Ile Trp Asp Glu Asp 30 35 40	509

gta Val 90	aaa Lys	gag Glu	aag Lys	gcc Ala	cat His 95	tca Ser	gga Gly	tcc Ser	cac His	agc Ser 100	gga Gly	ggt Gly	ggt Gly	cta Leu	gag Glu 105	701
					ctt Leu											749
aag Lys	gca Ala	ggc Gly	aaa Lys 125	gly ggg	gct Ala	agg Arg	gtg Val	ggt Gly 130	acc Thr	atc Ile	tct Ser	Gly 999	aac Asn 135	agg Arg	acc Thr	797
					ccc Pro											845
ccc Pro	acc Thr 155	agg Arg	agt Ser	gga Gly	tct Ser	agg Arg 160	gcc Ala	ggg Gly	ggc Gly	agg Arg	gca Ala 165	agt Ser	gga Gly	aaa Lys	tcc Ser	893
					agt Ser 175											941
cct Pro	gtc Val	cgg Arg	aga Arg	ggc Gly 190	aag Lys	ttc Phe	aac Asn	ttt Phe	cct Pro 195	tat Tyr	aaa Lys	att Ile	gat Asp	gat Asp 200	att Ile	989
					ctc Leu											1037
					caa Gln											1085
					aac Asn											1133
					ctg Leu 255											1181
					ctt Leu											1229
ggc	220	att	aad	acq	tac	atc	agt	caa	gtg	tgt	gat	gac	acc	atg	gtc	1277
GIY	Lys	Ile	Lys 285	Thr	Tyr	Ile	Ser	Gln 290	Val	Cys	Asp	Asp	Thr 295	Met	Val	

aac atg act gtg act aat cat tac caa cat ttg ctt tcc tat tct ttt Asn Met Thr Val Thr Asn His Tyr Gln His Leu Leu Ser Tyr Ser Phe 315 320 325	1373
cca gac ttt ttt gct ttg tta ttc ctg gga aat cac ttc acc aag ata Pro Asp Phe Phe Ala Leu Leu Phe Leu Gly Asn His Phe Thr Lys Ile 330 335 340 345	1421
cag att atg aaa cta att ata aac ttt act gaa aat cca gcc atg aca Gln Ile Met Lys Leu Ile Ile Asn Phe Thr Glu Asn Pro Ala Met Thr 350 355 360	1469
aga gag ctg gtc agt tgt aaa gta cca tca gaa ttg att tcc ctc ttt Arg Glu Leu Val Ser Cys Lys Val Pro Ser Glu Leu Ile Ser Leu Phe 365 370 375	1517
aat aaa gaa tgg gat aga gag att ctt ctt aat atc ctt acc cta ttt Asn Lys Glu Trp Asp Arg Glu Ile Leu Leu Asn Ile Leu Thr Leu Phe 380 385 390	1565
gag aat ata aat gac aac ata aaa aat gaa ggg ctc gca tca tcc agg Glu Asn Ile Asn Asp Asn Ile Lys Asn Glu Gly Leu Ala Ser Ser Arg 395 400 405	1613
aaa gaa ttc agc aga agt tca ctt ttt ttc tta ttc aaa gag tct gga Lys Glu Phe Ser Arg Ser Ser Leu Phe Phe Leu Phe Lys Glu Ser Gly 410 415 420 425	1661
gtt tgt gtt aag aaa atc aaa gca cta gca aat cac aat gat ctg gtg Val Cys Val Lys Lys Ile Lys Ala Leu Ala Asn His Asn Asp Leu Val 430 435 440	1709
gtg aaa gta aaa gtc ctg aaa gta tta acc aaa ctc taa tttggagtct Val Lys Val Leu Lys Val Leu Thr Lys Leu * 445 450	1758
gtcccaaaca atattgagat atttgcagtt ggtacgatgt gatttgtaaa ttctttgttt	1818
ttcattgtgc gtatatggta aagagatctt ttcagctgct attttggaat aatgactatc	1878
atatatcata acagtgactg atgttggttg taatggttgg gtttaggatg aaccatttta	1938
aggatgccaa atgaaatatt agtatttgta cacagaaaga atttattgat ttgatcttat	1998
tacctagatt gagatttttt aatctttcct ctacctaaac tgacaatgaa ttggttatac	2058
atcatgcata agctacactt ttatattagt ttatatttgt tattctaaga cttgtgtttc	2118
atcaataaag ttgtgtttta agcagcagaa aaaaaaaaaa	2162

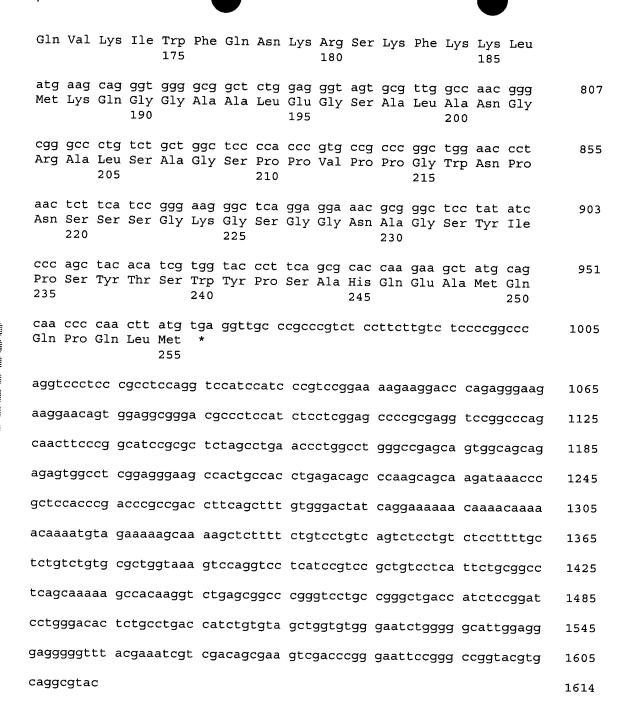
<210> 9 <211> 1614

<212> DNA

<213> Homo sapiens

<220>
<221> CDS
<222> (202)..(969)

<400> 9 aagctggtac gcctgcaggt accggtccgg aattcccggg tcgacgattt cgtcaaagag 60 acaaactcca ttttcttatg aatggaaagt gaaaacccct gttccgctta aattgggttc 120 180 cttcctqtcc tqaqaaacat agagaccccc aaaagggaag cagaggagag aaagtcccac acccagaccc cgcgagaaga g atg acc atg acc atg cca gaa agt ctc 231 Met Thr Met Thr Met Pro Glu Ser Leu aac agc ccc gtg tcg ggc aag gcg gtg ttt atg gag ttt ggg ccg ccc 279 Asn Ser Pro Val Ser Gly Lys Ala Val Phe Met Glu Phe Gly Pro Pro 15 327 aac cag caa atg tot cot toe occ atg toe cac ggg cac tac toe atg Asn Gln Gln Met Ser Pro Ser Pro Met Ser His Gly His Tyr Ser Met 30 375 cac tqt tta cac tcg gcg ggc cat tcg cag ccc gac ggc gcc tac agc His Cys Leu His Ser Ala Gly His Ser Gln Pro Asp Gly Ala Tyr Ser 45 423 tea gee teg tee tte tee ega eeg etg gge tae eec tae gte aac teg Ser Ala Ser Ser Phe Ser Arg Pro Leu Gly Tyr Pro Tyr Val Asn Ser gtc agc agc cac gca tcc agc ccc tac atc agt tcg gtg cag tcc tac 471 Val Ser Ser His Ala Ser Ser Pro Tyr Ile Ser Ser Val Gln Ser Tyr 519 ccq qqc aqc qcc agc ctc gcc cag agc cgc ctg gag gac cca ggg gcg Pro Gly Ser Ala Ser Leu Ala Gln Ser Arg Leu Glu Asp Pro Gly Ala 100 95 567 gac tog gag aag agc acg gtg gaa ggc ggt gaa gtg cgc ttc aat Asp Ser Glu Lys Ser Thr Val Val Glu Gly Glu Val Arg Phe Asn 115 110 ggc aag gga aaa aag atc cgt aaa ccc agg acg att tat tcc agt ttg 615 Gly Lys Gly Lys Lys Ile Arg Lys Pro Arg Thr Ile Tyr Ser Ser Leu 125 cag ttg cag gct ttg aac cgg agg ttc cag caa act cag tac cta gct 663 Gln Leu Gln Ala Leu Asn Arg Arg Phe Gln Gln Thr Gln Tyr Leu Ala 140 145 711 ctg ccg gag agg gcg gag ctc gcg gcc tct ttg gga ctc aca cag act Leu Pro Glu Arg Ala Glu Leu Ala Ala Ser Leu Gly Leu Thr Gln Thr 170 155 160 759 cag gtc aag atc tgg ttc caa aac aag cga tcc aag ttc aag aag ctg



```
<210> 10
```

<211> 4351

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (154)..(3240)

<400> 10 ccctctaatc tgatcactat agggaatttg gccctcgagc cgaagaattc ggcacgaggt	60
ctggctggga gcagaaggca gcctcggtct ctgggcggcg gcggcggccc actctgccct	120
ggccgcgctg tgtggtgacc gcaggcccga gac atg agg gcg gcc cgt gct ctg Met Arg Ala Ala Arg Ala Leu 1 5	174
ctg ccc ctg ctg ctg cag gcc tgc tgg aca gcc gcg cag gat gag ccg Leu Pro Leu Leu Gln Ala Cys Trp Thr Ala Ala Gln Asp Glu Pro 10 15 20	222
gag acc ccg agg gcc gtg gcc ttc cag gac tgc ccc gtg gac ctg ttc Glu Thr Pro Arg Ala Val Ala Phe Gln Asp Cys Pro Val Asp Leu Phe 25 30 35	270
ttt gtg ctg gac acc tct gag agc gtg gcc ctg agg ctg aag ccc tac Phe Val Leu Asp Thr Ser Glu Ser Val Ala Leu Arg Leu Lys Pro Tyr 40 45 50 55	318
ggg gcc ctc gtg gac aaa gtc aag tcc ttc acc aag cgc ttc atc gac Gly Ala Leu Val Asp Lys Val Lys Ser Phe Thr Lys Arg Phe Ile Asp 60 65 70	366
aac ctg agg gac agg tac tac cgc tgt gac cga aac ctg gtg tgg aac Asn Leu Arg Asp Arg Tyr Tyr Arg Cys Asp Arg Asn Leu Val Trp Asn 75 80 85	414
gca ggc gcg ctg cac tac agt gac gag gtg gag atc atc caa ggc ctc Ala Gly Ala Leu His Tyr Ser Asp Glu Val Glu Ile Ile Gln Gly Leu 90 95 100	462
acg cgc atg cct ggc ggc cgc gac gca ctc aaa agc agc gtg gac gcg Thr Arg Met Pro Gly Gly Arg Asp Ala Leu Lys Ser Ser Val Asp Ala 105 110	510
gtc aag tac ttt ggg aag ggc acc tac acc gac tgc gct atc aag aag Val Lys Tyr Phe Gly Lys Gly Thr Tyr Thr Asp Cys Ala Ile Lys Lys 120 125 130 135	558
ggg ctg gag cag ctc ctc gtg ggg ggc tcc cac ctg aag gag aat aag Gly Leu Glu Gln Leu Leu Val Gly Gly Ser His Leu Lys Glu Asn Lys 140 145 150	606
tac ctg att gtg gtg acc gac ggg cac ccc ctg gag ggc tac aag gaa Tyr Leu Ile Val Val Thr Asp Gly His Pro Leu Glu Gly Tyr Lys Glu 155 160 165	654
ccc tgt ggg ggg ctg gag gat gct gtg aac gag gcc aag cac ctg ggc Pro Cys Gly Gly Leu Glu Asp Ala Val Asn Glu Ala Lys His Leu Gly 170 175 180	702
gtc aaa gtc ttc tcg gtg gcc atc aca ccc gac cac ctg gag ccg cgt Val Lys Val Phe Ser Val Ala Ile Thr Pro Asp His Leu Glu Pro Arg 185 190 195	750

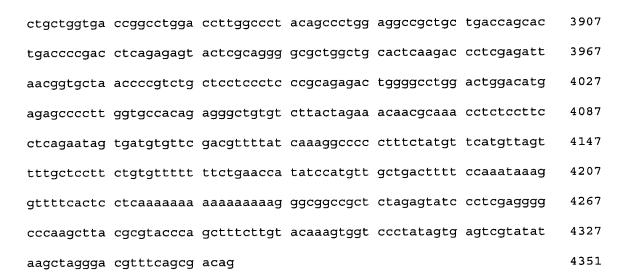
ctg Leu 200	agc Ser	atc Ile	atc Ile	gcc Ala	acg Thr 205	gac Asp	cac His	acg Thr	tac Tyr	cgg Arg 210	cgc Arg	aac Asn	ttc Phe	acg Thr	gcg Ala 215	798
			ggc													846
atc Ile	gac Asp	acc Thr	atc Ile 235	gtg Val	gac Asp	atg Met	atc Ile	aaa Lys 240	aat Asn	aac Asn	gtg Val	gag Glu	caa Gln 245	gtg Val	tgc Cys	894
			gaa Glu													942
gac Asp	ccc Pro 265	ggc Gly	ttt Phe	gag Glu	gga Gly	gaa Glu 270	cga Arg	ggc Gly	aag Lys	ccg Pro	999 Gly 275	ctc Leu	cca Pro	gga Gly	gag Glu	990
			gcc Ala													1038
GJÀ aaa	tac Tyr	cag Gln	gga Gly	atg Met 300	aag Lys	gga Gly	gaa Glu	aaa Lys	305 305	agc Ser	cgt Arg	Gly 999	gag Glu	aag Lys 310	ggc Gly	1086
			ccc Pro 315													1134
			gac Asp													1182
			ggc Gly													1230
ccc Pro 360	aag Lys	gga Gly	gac Asp	ccc Pro	ggc Gly 365	gcc Ala	ttt Phe	gga Gly	ctg Leu	aaa Lys 370	gga Gly	gaa Glu	aag Lys	ggc Gly	gag Glu 375	1278
cct Pro	gga Gly	gct Ala	gac Asp	999 380	gag Glu	gcg Ala	gly 999	aga Arg	cca Pro 385	ggg Gly	agc Ser	tcg Ser	gga Gly	cca Pro 390	tct Ser	1326
			ggc Gly 395													1374
			gac Asp													1422
cgg	ggt	ggc	cct	gga	gag	aga	gga	cca	cgg	ggg	acc	cca	ggc	acg	cgg	1470

Ar	g Gly 425		Pro	Gly	Glu	Arg 430	Gly	Pro	Arg	Gly	Thr 435	Pro	Gly	Thr	Arg	
	a cca y Pro 0															1518
	a gaa g Glu															1566
	c gga e Gly															1614
99 ¹	t gcc y Ala	aga Arg 490	gga Gly	gcc Ala	cca Pro	gga Gly	cct Pro 495	gcc Ala	gga Gly	ccc Pro	cct Pro	gga Gly 500	gac Asp	ccg Pro	Gly 999	1662
	g atg 1 Met 505															1710
	c ttc y Phe															1758
	a aac e Asn															1806
	ggg Gly															1854
	a gga s Gly															1902
gga Gl _y	a cac His 585	caa Gln	gga Gly	ccg Pro	cct Pro	999 Gly 590	ccg Pro	gac Asp	gaa Glu	tgc Cys	gag Glu 595	att Ile	ttg Leu	gac Asp	atc Ile	1950
	atg Met															1998
cto	ctg Leu	ttc Phe	gtg Val	ctg Leu 620	gac Asp	agc Ser	tca Ser	gag Glu	agc Ser 625	att Ile	ggc Gly	ctg Leu	cag Gln	aac Asn 630	ttc Phe	2046
gag	g att Ile	gcc Ala	aag Lys 635	gac Asp	ttc Phe	gtc Val	gtc Val	aag Lys 640	gtc Val	atc Ile	gac Asp	cgg Arg	ctg Leu 645	agc Ser	cgg Arg	2094
gac Asp	gag Glu	ctg Leu	gtc Val	aag Lys	ttc Phe	gag Glu	cca Pro	gly ggg	cag Gln	tcg Ser	tac Tyr	gcg Ala	ggt Gly	gtg Val	gtg Val	2142

650

660 cag tac agc cac agc cag atg cag gag cac gtg agc ctg cgc agc ccc 2190 Gln Tyr Ser His Ser Gln Met Gln Glu His Val Ser Leu Arg Ser Pro 665 670 age ate egg aac gtg cag gag ete aaq qaa qee ate aaq age etq caq 2238 Ser Ile Arg Asn Val Gln Glu Leu Lys Glu Ala Ile Lys Ser Leu Gln 685 tgg atg gcg ggc ggc acc ttc acg ggg gag gcc ctg cag tac acg cgg 2286 Trp Met Ala Gly Gly Thr Phe Thr Gly Glu Ala Leu Gln Tyr Thr Arg gac cag ctg ctg ccc ccc agc ccg aac aac cqc atc qcc ctq qtc atc 2334 Asp Gln Leu Leu Pro Pro Ser Pro Asn Asn Arg Ile Ala Leu Val Ile act gac ggg cgc tca gac act cag agg gac acc aca ccg ctc aac gtg 2382 Thr Asp Gly Arg Ser Asp Thr Gln Arg Asp Thr Thr Pro Leu Asn Val 730 ctc tgc agc ccc ggc atc cag gtg gtc tcc gtg ggc atc aaa gac gtg 2430 Leu Cys Ser Pro Gly Ile Gln Val Val Ser Val Gly Ile Lys Asp Val 745 750 ttt gac ttc atc cca ggc tca gac cag ctc aat gtc att tct tqc caa 2478 Phe Asp Phe Ile Pro Gly Ser Asp Gln Leu Asn Val Ile Ser Cys Gln ggc ctg gca cca tcc cag ggc cgg ccc ggc ctc tcg ctg gtc aag gaq 2526 Gly Leu Ala Pro Ser Gln Gly Arg Pro Gly Leu Ser Leu Val Lys Glu aac tat gca gag ctg ctg gag gat gcc ttc ctg aag aat gtc acc gcc 2574 Asn Tyr Ala Glu Leu Leu Glu Asp Ala Phe Leu Lys Asn Val Thr Ala 795 805 cag atc tgc ata gac aag aag tgt cca gat tac acc tgc ccc atc acg 2622 Gln Ile Cys Ile Asp Lys Lys Cys Pro Asp Tyr Thr Cys Pro Ile Thr 810 ttc tcc tcc ccg gct gac atc acc atc ctg ctg gac ggc tcc gcc agc 2670 Phe Ser Ser Pro Ala Asp Ile Thr Ile Leu Leu Asp Gly Ser Ala Ser 825 830 gtg ggc agc cac aac ttt gac acc acc aag cgc ttc gcc aag cgc ctq 2718 Val Gly Ser His Asn Phe Asp Thr Thr Lys Arg Phe Ala Lys Arg Leu 840 855 2766 Ala Glu Arg Phe Leu Thr Ala Gly Arg Thr Asp Pro Ala His Asp Val 860 cgg gtg gcg gtg gtg cag tac agc ggc acg ggc cag cag cgc cca gaq 2814 Arg Val Ala Val Val Gln Tyr Ser Gly Thr Gly Gln Gln Arg Pro Glu 875 880

cgg Arg	gcg Ala	tcg Ser 890	ctg Leu	cag Gln	ttc Phe	ctg Leu	cag Gln 895	aac Asn	tac Tyr	acg Thr	gcc Ala	ctg Leu 900	gcc Ala	agt Ser	gcc Ala	2862
gtc Val	gat Asp 905	gcc Ala	atg Met	gac Asp	ttt Phe	atc Ile 910	aac Asn	gac Asp	gcc Ala	acc Thr	gac Asp 915	gtc Val	aac Asn	gat Asp	gcc Ala	2910
ctg Leu 920	ggc Gly	tat Tyr	gtg Val	acc Thr	cgc Arg 925	ttc Phe	tac Tyr	cgc Arg	gag Glu	gcc Ala 930	tcg Ser	tcc Ser	ggc Gly	gct Ala	gcc Ala 935	2958
aag Lys	aag Lys	agg Arg	ctg Leu	ctg Leu 940	ctc Leu	ttc Phe	tca Ser	gat Asp	ggc Gly 945	aac Asn	tcg Ser	cag Gln	ggc Gly	gcc Ala 950	acg Thr	3006
ccc Pro	gct Ala	gcc Ala	atc Ile 955	gag Glu	aag Lys	gcc Ala	gtg Val	cag Gln 960	gaa Glu	gcc Ala	cag Gln	cgg Arg	gca Ala 965	ggc Gly	atc Ile	3054
gag Glu	atc Ile	ttc Phe 970	gtg Val	gtg Val	gtc Val	gtg Val	ggc Gly 975	cgc Arg	cag Gln	gtg Val	aat Asn	gag Glu 980	ccc Pro	cac His	atc Ile	3102
cgc Arg	gtc Val 985	ctg Leu	gtc Val	acc Thr	ggc Gly	aag Lys 990	acg Thr	gcc Ala	gag Glu	tac Tyr	gac Asp 995	gtg Val	gcc Ala	tac Tyr	ggc Gly	3150
gag Glu 1000	agc Ser	cac His	ctg Leu	Phe	cgt Arg 1005	gtc Val	ccc Pro	agc Ser	Tyr	cag Gln 1010	gcc Ala	ctg Leu	ctc Leu	cgc Arg	ggt Gly 1015	3198
gtc Val	ttc Phe	cac His	Gln	aca Thr 1020	gtc Val	tcc Ser	agg Arg	Lys	gtg Val 1025	Ala	ctg Leu	ggc Gly	tag *	ccc	accc	3247
tgc	acgc	cgg	cacc	aaac	cc t	gtcc	tccc	a cc	cctc	ccca	ctc	atca	cta	aaca	gagaaa	3307
agc	ttgg	aaa	gcca	ggac	ac a	acgc	tgct	g cc	tgct	ttgt	gca	gggt	cct	ccgg	ggctca	3367
gcc	ctga	gtt	ggca	tcac	ct g	cgca	gggc	c ct	ctgg	ggct	cag	ccct	gag	ctag	tgtcac	3427
ctg	caca	999	ccct	ctgg	gg c	tcag	ccct	g ag	ctgg	cgtc	acc	tgtg	cag	ggcc	ctctgg	3487
ggd	tcag	ccc	tgag	ctgg	cc t	cacc	tggg	t to	ссса	.cccc	999	ctct	cct	gccc	tgccct	3547
cct	gccc	gcc	ctcc	ctcc	tg c	ctgc	gcag	c to	ctto	ccta	ggc	acct	ctg	tgct	gcgtcc	3607
cac	cago	ctg	agca	agac	gc c	ctct	cggg	g co	tgtg	ccgc	act	agco	tcc	ctct	cctctg	3667
tcc	ccat	agc	tggt	tttt	.cc c	acca	atco	t ca	ccta	acag	, tta	cttt	aca	atta	aactca	3727
aag	gcaag	gctc	ttct	cctc	ag c	ttgg	ıggca	ig co	attg	gcct	ctg	gtete	gtt	ttgg	gaaacc	3787
aag	gtca	ıgga	ggco	gttg	rca g	jacat	aaat	c to	ggcg	gacto	ggo	cccc	gtct	ccts	gagggtc	3847



<210> 11 <211> 1622 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (864)..(1424)

<400> 11 60 taagettgeg geegetgaac tacagetetg egeetgeeca ggeggeegea egeteagggg cgtggcatgg gtgggtcgtg agttgggcgg ggcccacagg gcgtgcgcga cgcagcggcg 120 cggcgcgagg cgtaaggggc gtggcgccag tgggcgtggc gtggcgcagt gcgaagggac 180 geggtgegea tgegegtgag ggetteeaeg ggtgggtggt ategaggeet gtegggteag 240 ggcggttcgc gggtgctgtc agagctgggc cggggcccct aggcaggccc agacatgtcc 300 gtccttgtaa gttaaaagct tccatgggag ccttccttcc taatcaagat gcaaatagta 360 cggtattccg aacagacact aaaaatagct gtcatctcaa agaatccagt gcttgtgtca 420 cagtatgaga aagtagatgc tggggaacag cgtttaatga atgaggcatg caagccagcc 480 agtgatetet ttggaeettg cattetecat cagattggat caceteceae cetgaggeee 540 cccaagactt tgaacagttc ttcagtcatc cttacagaaa gataccctct ccagacaaac 600 660 gcagtattta tatacggtcc attggatctc tatgaagcac cagaattatc agtgaagaat 720 atattaaatg gctcacgggc tactgtaaag catatttcta tcgcttgaga gtaaaactcc 780 tagaaccagt teetgtttet gtaacaagat gtteetttag agteaatgag aacacacaca

acctacaaat tcatgcaggg gacatcctga agttcttgaa aaagaagaaa cctgaagatg 8											
ccttctgtgt tgtgggaata aca atg att gat ctt tac cca aga gac tcg Met Ile Asp Leu Tyr Pro Arg Asp Ser 1 5	890										
tgg aat ttt gtc ttt gga cag gcc tct ttg aca gat ggt gtg ggg ata Trp Asn Phe Val Phe Gly Gln Ala Ser Leu Thr Asp Gly Val Gly Ile 10 15 20 25	938										
ttc agc ttt gcc agg tat ggc agt gat ttt tat agc atg cac tat aaa Phe Ser Phe Ala Arg Tyr Gly Ser Asp Phe Tyr Ser Met His Tyr Lys 30 35 40	986										
ggc aaa gtg aag aag ctc aag aaa aca tct tca agt gac tat tca att Gly Lys Val Lys Lys Leu Lys Lys Thr Ser Ser Ser Asp Tyr Ser Ile 45 50 55	1034										
ttc gac aac tat tat att cca gaa ata act agt gtt tta cta ctt cga Phe Asp Asn Tyr Tyr Ile Pro Glu Ile Thr Ser Val Leu Leu Leu Arg 60 65 70	1082										
tcc tgt aag act tta acc cat gag atc gga cac ata ttt gga ctg cga Ser Cys Lys Thr Leu Thr His Glu Ile Gly His Ile Phe Gly Leu Arg 75 80 85	1130										
cac tgc cag tgg ctt gca tgc ctc atg caa ggc tcc aac cac ttg gaa His Cys Gln Trp Leu Ala Cys Leu Met Gln Gly Ser Asn His Leu Glu 90 95 100 105	1178										
gaa gct gac cgg cgc cct cta aac ctt tgc cct atc tgt ttg cac aag Glu Ala Asp Arg Arg Pro Leu Asn Leu Cys Pro Ile Cys Leu His Lys 110 115 120	1226										
ttg cag tgt gct gtt ggc ttc agc att gta gaa aga tac aaa gca ctg Leu Gln Cys Ala Val Gly Phe Ser Ile Val Glu Arg Tyr Lys Ala Leu 125 130 135	1274										
gtg agg tgg att gat gat gaa tct tct gac aca cct gga gca act cca Val Arg Trp Ile Asp Asp Glu Ser Ser Asp Thr Pro Gly Ala Thr Pro 140 145 150	1322										
gaa cac agt cac gag gat aat ggg aat tta ccg aaa ccc gtg gaa gcc Glu His Ser His Glu Asp Asn Gly Asn Leu Pro Lys Pro Val Glu Ala 155 160 165	1370										
ttt aag gaa tgg aaa gag tgg ata ata aaa tgc ctg gct gtt ctc caa Phe Lys Glu Trp Lys Glu Trp Ile Ile Lys Cys Leu Ala Val Leu Gln 170 175 180 185	1418										
aaa tga ggaccttcaa ataggagtga ttgaaataaa taactacttg catgttatgc Lys *	1474										
tttcatttgg gtggaatact tcattggaat aaactactga tcttgtgctg tgtcaaagta	1534										
acagactaga accttctttc aagtacctga attgaaatga aactcatttt gaataataaa	1594										

<210> 12 <211> 882 <212> DNA <213> Homo sapiens	
<220> <221> CDS <222> (147)(284)	
<400> 12	
gagccgagat cacgccactg cactccagcc tgggcaacaa gagcaaaact ctgtctcaat	60
aaaaaaaag gaaaacccac agccaacatt atcactagtg gtaacaggga aaatgtcccc 1	20
tttgaccccc accccaagta cagaac atg caa gga cgc ctg ttc tca ctg cat Met Gln Gly Arg Leu Phe Ser Leu His 1 5	.73
gtc ttc tgt gtt gta ctg aag ctc cta gct ggt gca gtg agg caa gac Val Phe Cys Val Val Leu Lys Leu Leu Ala Gly Ala Val Arg Gln Asp 10 15 20 25	21
aaa gaa agg aaa tat agg tgc act ggg aag gaa gaa gaa aca ctg cct Lys Glu Arg Lys Tyr Arg Cys Thr Gly Lys Glu Glu Glu Thr Leu Pro 30 35 40	69
tta ttc ttt agg tga catgattgtg tgcttttaaa ataataaagg aatcaacaga 3 Leu Phe Phe Arg * 45	24
aaagttgctt aacctaatga atgagtttat caaagtcaca agatacaagg tcagtataca 3	84
aaaatcagtt ggatttctac atggtagaaa caactgtaca tggaaaaatg tttaatagtg 4	44
taagatatgt acattggaaa ctatgaaaga gtgtaaaaaa taaagaagtg aaataaat	04
agaagatacc accttgatgg atggaagcct taaggtaaag atgctcattc tccccacact 5	64
gacctgtaca ttccccacag gcctaatcaa aacccccaca ggcttctgtg tagaaattga 6	24
catgctgatc ctgaaattta tatgaaaatg caaagagtct ggaataacca aaataatttt 6	84
gtaaaagaac aaagaagact totactacct ggttataaga cttctctgaa gcacagaagt 74	44
caaggcagtg tggtggtggc ataagtaatg taaatcatcc aggtgtggtg gctcaagcca 80	04
gtcatccagc actttgggag gctgaggcag gagaatcgct agagcccagg agttggaaac 80	64
cagcctgggc agcatagc 88	82

<210> 13

<221> CDS

<222> (173)..(1909)

	<2	11> 4 12> 1	DNA													
	<2	13> 1	HOMO	sap:	iens											
		20>														
		21> (\											
	<2	22>	(58)	(3.	36)											
	<4	00>	13													
													-	actg	_	57
														cag		105
Met 1	Tyr	тте	гаг	Pro 5	ser	Thr	Arg	га	ser 10	GIY	Tyr	ser	Pro	Gln 15	GIN	
1				,					10					15		
gta	gct	gtg	atc	cac	tgc	aaa	gga	cat	caa	aaa	gaa	aac	acg	gcc	gtt	153
Val	Ala	Val		His	Cys	Lys	Gly		Gln	Lys	Glu	Asn		Ala	Val	
			20					25					30			
acc	cat	agt	aac	caq	aaa	act	gat	tca	gca	act	caq	atc	act	gcc	aga	201
														Ala		
		35					40					45				
	.											.				240
		_	_				_	_			_			cca Pro	_	249
Бец	50	vai	1111	FIU	FIO	55	пеп	пец	PIO	1111	60	per	PIIC	PIO	GIII	
														aaa		297
	Asp	Leu	Pro	Asp		Pro	Val	Tyr	Ser		Thr	Thr	Glu	Lys		
65					70					75					80	
gct	tca	gat	ctc	aqa	qcc	aat	aaa	aat	caq	qaa	aqt	taq	taga	attct	tc	346
								Asn				*				
				85					90							
atas	ata	-~~	. a t a t	-++		-a+a	at			·++ ~	aata				+	106
ctga	acte	tgg a	acci	LCai	La CC	octga	adcu	L ada	accaç	jila	CCL	icagi	.CL a	accac	ccatt	406
taag	gaaga	agc a	aaagt	taco	ct ca	agcto	cata	gga	aggg							442
	<2	10> 1	L 4													
	<23	11> 2	2058													
		12> I														
	<21	13> F	Omo	sapi	lens											
	<22	20>														

60

120

taagcttgcg gccgcccggc ggtgtagatg gtacgcttgc ttgtacgcag caggggaggc

tetgaettge aetggetggt ggteegette eggggagage geeeagtgee caeaggtgge

aggagcct	gt ccago	ggactc t	gcatt	gctg	g ctg	gcaga	agag	ggat	gtad	etc g	aa	atg Met 1	1	175
tgg ccc a													2	223
aca ggg (Thr Gly I													2	271
cct gtt q Pro Val (35													3	319
ggg tac o Gly Tyr 1 50													3	367
ctc tgc a													4	415
ggc act o													4	463
gag cct (Glu Pro													Ē	511
cgg ccg (Arg Pro 1													Ę	559
gcc atc q Ala Ile (130													6	607
aga atg Arg Met													•	655
atg gac Met Asp													•	703
gca cac Ala His													•	751
ctc atc Leu Ile 195			_	_	_	_	_		_	_	-	_	•	799

·	
gtg ccc att tat gac ttc acc acg cac agc cgg aag aag gac tgg aaa Val Pro Ile Tyr Asp Phe Thr Thr His Ser Arg Lys Lys Asp Trp Lys 210 225	847
aca ctg tat ggt gca aac gtc atc atc ttt gag ggc atc atg gcc ttt Thr Leu Tyr Gly Ala Asn Val Ile Ile Phe Glu Gly Ile Met Ala Phe 230 230 240	895
gct gac aag aca ctg ttg gag ctc ctg gac atg aag atc ttt gtg gac Ala Asp Lys Thr Leu Leu Glu Leu Leu Asp Met Lys Ile Phe Val Asp 255	943
aca gac tcc gac atc cgc ctg gta cgg cgg ctg cgc cgg gac atc agt Thr Asp Ser Asp Ile Arg Leu Val Arg Arg Leu Arg Arg Asp Ile Ser 260 265 270	991
gag cgc ggc cgg gac atc gag ggt gtc atc aag cag tac aac aag ttt Glu Arg Gly Arg Asp Ile Glu Gly Val Ile Lys Gln Tyr Asn Lys Phe 275 280 285	1039
gtc aag ccc tcc ttc gac cag tac atc cag ccc acc atg cgc ctg gca Val Lys Pro Ser Phe Asp Gln Tyr Ile Gln Pro Thr Met Arg Leu Ala 305	1087
gac atc gtg gtc ccc aga ggg agc ggc aac acg gtg gcc atc gac ctg Asp Ile Val Val Pro Arg Gly Ser Gly Asn Thr Val Ala Ile Asp Leu 310 310	1135
att gtg cag cac gtg cac agc cag ctg gag gag cgt gaa ctc agc gtc Ile Val Gln His Val His Ser Gln Leu Glu Glu Arg Glu Leu Ser Val 325 326	1183
agg gct gcg ctg gcc tcg gca cac cag tgc cac ccg ctg ccc cgg acg Arg Ala Ala Leu Ala Ser Ala His Gln Cys His Pro Leu Pro Arg Thr 340	1231
ctg agc gtc ctg aag agc acg ccg cag gta cgg ggc atg cac acc atc Leu Ser Val Leu Lys Ser Thr Pro Gln Val Arg Gly Met His Thr Ile 355	1279
atc agg gac aag gag acc agt cgc gac gag ttc atc ttc tac tcc aag Ile Arg Asp Lys Glu Thr Ser Arg Asp Glu Phe Ile Phe Tyr Ser Lys 370 385	1327
aga ctg atg cgg ctg ctc atc gag cac gcg ctc tcc ttc ctg ccc ttt aga ctg atg cgg ctg ctc atc gag cac gcg ctc tcc ttc ctg ccc ttt Arg Leu Met Arg Leu Leu Ile Glu His Ala Leu Ser Phe Leu Pro Phe 390 395	1375
cag gac tgc gtc gta cag acc ccg cag ggg cag gac tat gcg ggc aag Gln Asp Cys Val Val Gln Thr Pro Gln Gly Gln Asp Tyr Ala Gly Lys 405	1423
tgc tat gcg ggg aag cag atc acc ggt gtg tcc att ctg cgc gcc ggt Cys Tyr Ala Gly Lys Gln Ile Thr Gly Val Ser Ile Leu Arg Ala Gly 420 425	1471
gaa acc atg gag ccc gcg ctg cgc gct gtg tgc aaa gac gtg cgc atc	1519

Glu Thr Met Glu Pro Ala Leu Arg Ala Val Cys Lys Asp Val Arg Ile 435 440 445	
ggc acc atc ctc atc cag acc aac cag ctt acc ggg gag ccc gag ctc Gly Thr Ile Leu Ile Gln Thr Asn Gln Leu Thr Gly Glu Pro Glu Leu 450 465 460 465	1567
cac tac ctg agg ctg ccc aag gac atc agc gat gac cac gtg atc ctc His Tyr Leu Arg Leu Pro Lys Asp Ile Ser Asp Asp His Val Ile Leu 470 475 480	1615
atg gac tgc acc gtg tcc acg ggc gcg gcc atg atg gca gtg cgc Met Asp Cys Thr Val Ser Thr Gly Ala Ala Ala Met Met Ala Val Arg 485 490 495	1663
gtg ctc ctg gac cac gac gtg cct gag gac aag atc ttt ttg ctg tcg Val Leu Leu Asp His Asp Val Pro Glu Asp Lys Ile Phe Leu Leu Ser 500 505 510	1711
ctg ctc atg gca gag atg ggc gtg cac tca gtg gcc tat gca ttt ccg Leu Leu Met Ala Glu Met Gly Val His Ser Val Ala Tyr Ala Phe Pro 515 520 525	1759
cga gtg aga atc atc acc acg gcg gtg gac aag cgg gtc aat gac ctt Arg Val Arg Ile Ile Thr Thr Ala Val Asp Lys Arg Val Asn Asp Leu 530 545	1807
ttc cgc atc atc cca ggc att ggg aac ttt ggc gac cgc tac ttt ggg Phe Arg Ile Ile Pro Gly Ile Gly Asn Phe Gly Asp Arg Tyr Phe Gly 550 555 560	1855
aca gac gcg gtc ccc gat ggc agt gac gag gag gaa gtg gcc tac acg Thr Asp Ala Val Pro Asp Gly Ser Asp Glu Glu Glu Val Ala Tyr Thr 565 570 575	1903
ggt tag ctgcccagtg agccatcccg tccccaccac cctcctcctg cctcctgacc Gly *	1959
caggactgct gaatacaaag atgttaattt ttaaaatgtt actagtataa tttattctat	2019
gcattttata aaataaataa agctttagaa aaaaaaaaa	2058

```
<210> 15
<211> 1705
<212> DNA
<213> Homo sapiens
<220>
<221> CDS
<222> (545)..(1672)
```

<400> 15
aaggateett aattttatta ateceeece eeeegaagge eeeeggeege etgtgeeeeg 60

cgg	tgcg	cgc	cccg	jetec	gg g	acct	gccg	c ca	ccgc	cgcc	ccg	ccct	cgg	cggc	acccac	120
acc	cagg	cgc	gccc	gcgc	gc g	cgcc	cggc	c cc	gtcc	ctgo	ctg	gaag	cac	agct	gaagat	180
ggc	gago	ccg	gcgc	ctcc	gg a	gcac	gccg	a gg	aggg	atgo	ccg	gcto	ctg	ccgc	cgagga	240
gca	ggcg	ccg	ccgt	cgcc	gc c	accg	cccc	a gg	cato	cccc	gca	gago	ggc	agca	gcagga	300
gga	ggaa	gcg	cagg	aago	tg g	ggcg	gcgg	a gg	gcgc	aaaa	ttg	cagg	tgg	agga	ggccgc	360
3 39	ccgg	gcg	gcgg	ccgc	gg t	aacc	tggc	t ga	tcgg	ggag	ccg	gtgc	tgt	ggct	gggctg	420
ccg	cgcc	gac	gagc	tgct	ga g	ctgg	aaga	g gc	cgct	gcgg	agc	ctgc	tcg	gctt	cgtcgc	480
tgc	caac	ctg	ctgt	tctg	gt t	cctt	gcat	t ga	ctcc	atgg	aga	gtat	atc	acct	gatttc	540
cgt	Me	g at t Il 1	a ct e Le	t gg u Gl	y Ar	t gt g Va 5	t at l Il	t at e Me	g ca t Gl	a at n Il 1	e Il	a aa e Ly	g ga s As	t at p Me	g gtt t Val 15	589
ttg Leu	tct Ser	aga Arg	aca Thr	aga Arg 20	ggt Gly	gca Ala	cag Gln	ttg Leu	tgg Trp 25	aga Arg	agc Ser	ctc Leu	agt Ser	gaa Glu 30	agc Ser	637
tgg Trp	gaa Glu	gtt Val	atc Ile 35	aat Asn	tcc Ser	aaa Lys	cca Pro	gat Asp 40	Glu	aga Arg	ccc Pro	agg Arg	ctc Leu 45	agc Ser	cac His	685
tgt Cys	att Ile	gca Ala 50	gaa Glu	tca Ser	tgg Trp	atg Met	aat Asn 55	ttc Phe	agc Ser	ata Ile	ttt Phe	ctt Leu 60	caa Gln	gaa Glu	atg Met	733
tct Ser	ctt Leu 65	ttt Phe	aaa Lys	cag Gln	cag Gln	agc Ser 70	cct Pro	ggc Gly	aag Lys	ttt Phe	tgt Cys 75	ctc Leu	ctg Leu	gtc Val	tgt Cys	781
agt Ser 80	gtg Val	tgc Cys	aca Thr	ttt Phe	ttt Phe 85	acg Thr	atc Ile	ttg Leu	gga Gly	agt Ser 90	tac Tyr	att Ile	cct Pro	ggg Gly	gtt Val 95	829
ata Ile	ctc Leu	agc Ser	tat Tyr	cta Leu 100	ctg Leu	tta Leu	ctg Leu	tgt Cys	gca Ala 105	ttt Phe	ttg Leu	tgt Cys	cca Pro	ttg Leu 110	ttt Phe	877
aaa Lys	tgt Cys	aat Asn	gat Asp 115	att Ile	gga Gly	caa Gln	aaa Lys	att Ile 120	tac Tyr	agc Ser	aaa Lys	att Ile	aag Lys 125	tca Ser	gtt Val	925
ctg Leu	ctg Leu	aaa Lys 130	ctg Leu	gat Asp	ttt Phe	gga Gly	att Ile 135	gga Gly	gaa Glu	tat Tyr	att Ile	aat Asn 140	cag Gln	aag Lys	aaa Lys	973
cgt Arg	gag Glu 145	aga Arg	tct Ser	gaa Glu	gca Ala	gat Asp 150	aaa Lys	gaa Glu	aaa Lys	agt Ser	cac His 155	aaa Lys	gat Asp	gac Asp	agt Ser	1021

						ctt Leu											1069
_						tct Ser	_		_	_			_				1117
						aac Asn											1165
_			_	_		gac Asp	_		_		_	_			_		1213
_			_			tct Ser 230		_			_				_		1261
						ggt Gly											1309
_	_	_	_	_		cac His	_		_			_			_		1357
_						ctg Leu		_	_					_	_		1405
_		_	_		_	gtt Val			_	_			_				1453
	_	_				gtg Val 310	_		_			_	-	_			1501
						gac Asp											1549
						gat Asp											1597
						gca Ala										:	1645
						ggc Gly		taa *	tcta	ag ga	aatca	igcti	gca	aacag	gagc		1697
aca	aaaa	ā															1705

<210> 16

	<2	11>	1914													
	<2	12>	DNA													
	<2	13>	Homo	sap	iens											
	-2	20>														
		21>	CDS													
	<2	22>	(154) (1593)										
	_															
qqqa		00> tat :		atga	ta a	cact	aqta	c ca	teca	acct	att	tagg	tga	cact	atagaa	60
555	333	•	33	J		J			J	J		55	- 5			
caag	gttt	gta	caaa	aaag	ca g	gctg	gtac	c gg	tccg	gaat	tcc	cggg	ata	tcgt	cgaccc	120
acgo	cgtc	cgg	tgca	ccgc	gt t	ctcg	cacg	c gt	Me						a cag l Gln	174
ctg	gtg	gtg	acc	ctg	ctc	act	gcc	acc	ctc	atg	cac	agg	ctg	gcg	cca	222
															Pro	
cac	tgc	tcc	ttc	gcg	cgc	tgg	ctg	ctc	tgt	aac	ggc	agt	ttg	ttc	cga	270
	Cys		Phe			Trp					Gly					
	25					30					35					
tac	aaq	cac	ccg	tct	qaq	qaq	qaq	ctt	caa	acc	cta	aca	aaa	aaq	cca	318
Tyr	Lys	His	Pro	Ser	Glu	Glu	Glu	Leu	Arg	Ala	Leu	Ala	Gly	Lys	Pro	
40					45					50					55	
agg	ccc	aga	ggc	agg	aaa	gag	caa	taa	acc	aat.	aac	ctt	agt.	gag	gag	366
			Gly													300
				60					65					70		
aaa	cca	cta	tct	ata	ddd	cga	gat	acc	cca	ttc	cag	cta	gag	acc	tac	414
			Ser													272
			75				_	80					85		-	
cc	ctc	aco	acc	ata	gat	acc	ata	atc	cta	כמכ	ttc	ttc	cta	asa	tac	462
			Thr													402
		90			_		95			_		100			-	
'ag	taa	+++	gtg	gac	+++	act	ata	tac	tca	aac	aac	ata	tac	ctc	ttc	510
			Val													310
	105					110		-		-	115		-			
ca	gag	acc	tac	tac	tac	ato	cta	aas	cca	acc	220	aaa	act	220	a++	558
			Tyr													338
L20			=	-	125			-		130	-				135	
ıct	ata	ttc	tgg	tac	cta	ctc	aco	ata	acc	ttc	tcc	ato	227	ato	ttc	606
			Trp													000
			_	140					145				-	150		

	ctg Leu	aca Thr	gto Va	L T	ca c hr <i>I</i> 55	egg Arg	ctg Leu	tac Tyr	ttc Phe	agc Ser 160	gc Al	с g a G	ag (gag Glu	Gly 999	ggt Gly 165	_	ag llu	cg Ar	c g	654
	tct Ser	gtc Val	tg Cy 17	s L	tc a eu :	acc Thr	ttt Phe	gcc Ala	ttc Phe 175	ctc Leu	tt Ph	.c c	tg eu	ctg Leu	ctg Leu 180	gco	c a	itg Iet	ct Le	g u	702
	gtg Val	caa Gln 185	Va	g g 1 V	tg (cgg Arg	gag Glu	gag Glu 190	acc Thr	ctc Leu	ga Gl	ig o Lu I	etg Leu	ggc Gly 195	ctg Leu	gaq	g c	cct Pro	gg Gl	t Y	750
	ctg Leu 200	gcc	ag Se	c a	itg Iet	acc Thr	cag Gln 205	aac Asn	tta Leu	gag	g CC		ctt Leu 210	ctg Leu	aag Lys	aa Ly	g (s (cag Gln	gg Gl 21	-	798
	tgg Trp	gad Ası	tg Tr	ıg g	gcg Ala	ctt Leu 220	cct Pro	gtg Val	gcc Ala	aag Lys	э ш t	tg (eu . 25	gct Ala	atc Ile	cgc Arg	gt y Va	g (gga Gly 230	ct Le	eu	846
	gca Ala	gte Va	g gt l Va	al (ggc Gly 235	tct Ser	gtg Val	ctg Leu	ggt Gly	gco Ala 24	3 P.	tc he	ctc Leu	acc Thr	tto Phe	c cc e Pr 24	_	ggc Gly	ct Le	tg eu	894
	cgg Arg	g ct g Le	u A	cc (la (cag Gln	acc Thr	cac	cgg Arg	gaq Asp 255) AI	a c a L	tg eu	acc Thr	atg Met	tc Se 26		ig Lu	gac Asp	a A	ga rg	942
	cco Pro	at Me 26	t L	tg eu	cag Gln	ttc Phe	cto Lev	cto Lei 27	g cad u His	c ac s Th	c a r S	igc Ser	ttc Phe	cto Lei 275	, ,,,	t co r Pi	cc	ctg Lev	j t 1 P	tc he	990
tant timi indi il	ate [1] 28	e Le	g t u T	gg rp	ctc Leu	tgg Trp	g aca Thi	с ГА	g cc s Pr	c at o Il	t c .e <i>P</i>	gca Ala	cgg Arg 290	, AS	e tt o Ph	c ct	tg eu	cac His	c c s G 2	ag In 195	1038
ž	cc Pr	g co o Pi	g t	tt	ggg ggg	gag Glu	ı Th	g cg r Ar	t tt g Ph	c to e Se	er 1	ctg Leu 305	cto	g to i Se	c ga r As	t t p S	ct er	gco Ala 31		tc Phe	1086
	ga As	c to p Se	ct g er G	31y 399	cgc Arg	, Le	c tg u Tr	g tt p Le	g ct u Le	u v	=g 9 al 1 20	gtg Val	cto	g tg ı Cy	c ct s Le		tg eu 25		g (ctg Leu	1134
	gc Al	g g .a V	al 7	acc Thr	cgg	g cc g Pr	с са о Ні	c ct s Le	g ca eu Gl 33	n A	cc la	tac Tyr	ct; Le	g tg u Cy	<u>.</u>	=g g =u A 40	ıcc Ma	aa Ly	g g	gcc Ala	1182
	cg Ar	g V	tg q al (gag Glu	caq Gl:	g ct n Le	g cg u Ar	g Ai	gg ga gg G]	ag g Lu A	ct la	ggc Gly	cg Ar	c at g Il 35		aa g lu <i>F</i>	gcc Ala	cg Ar	gt (gaa Glu	1230
	I.	cc c le G	ag ln	cag Gln	agg Arg	g gt g Va	ıl Va	cc cg al A:	ga gi rg Va	tc t al T	ac 'yr	tgo Cys	ta Ty 37		g a al T	cc 9	gt <u>c</u> Va]	g gt L Va	g al	agc Ser 375	1278

	ttg cag tac ctg acg ccg ctc atc ctc acc ctc aac tgc aca ctt ctg Leu Gln Tyr Leu Thr Pro Leu Ile Leu Thr Leu Asn Cys Thr Leu Leu 380 385 390	1326
	ctc aag acg ctg gga ggc tat tcc tgg ggc ctg ggc cca gct cct cta Leu Lys Thr Leu Gly Gly Tyr Ser Trp Gly Leu Gly Pro Ala Pro Leu 395	1374
	cta tcc ccc gac cca tcc tca gcc agc gct gcc ccc atc ggc tct ggg Leu Ser Pro Asp Pro Ser Ser Ala Ser Ala Ala Pro Ile Gly Ser Gly 410 415 420	1422
	gag gac gaa gtc cag cag act gca gcg cgg att gcc ggg gct ctg ggt Glu Asp Glu Val Gln Gln Thr Ala Ala Arg Ile Ala Gly Ala Leu Gly 425 430 435	1470
	ggc ctg ctt act ccc ctc ttc ctc cgt ggc gtc ctg gcc tac ctc atc Gly Leu Leu Thr Pro Leu Phe Leu Arg Gly Val Leu Ala Tyr Leu Ile 450 455	1518
Hoth Hoth II	tgg tgg acg gct gcc tgc cag ctg ctc gcc agc ctt ttc ggc ctc tac Trp Trp Thr Ala Ala Cys Gln Leu Leu Ala Ser Leu Phe Gly Leu Tyr 460 465 470	1566
	ttc cac cag cac ttg gca ggc tcc tag ctgcc tgcagaccct cctggggccc Phe His Gln His Leu Ala Gly Ser * 475 480	1618
7 (m)	tgaggtetgt teetggggea gegggaeaet ageetgeeee etetgtttge geeeeegtgt	1678
	ccccagctgc aaggtggggc cggactcccc ggcgttccct tcaccacagt gcctgacccg	1738
	cggcccccct tggacgccga gtttctgcct cagaactgtc tctcctgggc ccagcagcat	1798
:41	gagggtcccg aggccattgt ctccgaagcg tatgtgccag gtttgagtgg cgagggtgat	1858
H H W H	gctggctgct cttctgaaca aataaaggag catgccgatt tttacaaaaa aaaaaa	1914

```
<210> 17
    <211> 859
    <212> DNA
    <213> Homo sapiens
    <220>
    <221> CDS
     <222> (61)..(795)
atttggccct cgaggccaag aattcggcac gaggaaggag gtcccccacg gcccttcagg
                                                                      60
atg aaa gct gcg gtg ctg acc ttg gcc gtg ctc ttc ctg acg ggg agc
                                                                      108
Met Lys Ala Ala Val Leu Thr Leu Ala Val Leu Phe Leu Thr Gly Ser
                                     10
                  5
  1
```

cag Glr	gct Ala	cgg Arg	cat His 20	Phe	tgg Trp	cag Gln	caa Gln	gat Asp 25	Glu	. ccc . Pro	ccc Pro	cag Gln	ago Ser 30	Pro	tgg Trp	156
gat Asp	cga Arg	gtg Val 35	Lys	gac Asp	ctg Leu	gcc Ala	act Thr 40	Val	tac Tyr	gtg Val	gat Asp	gtg Val 45	Leu	aaa Lys	gac Asp	204
agc Ser	ggc Gly 50	Lys	gac Asp	agc Ser	gtg Val	acc Thr 55	tcc Ser	acc Thr	ttc Phe	agc Ser	aag Lys 60	Leu	cgc Arg	gaa Glu	cag Gln	252
ctc Leu 65	Gly	cct Pro	gtg Val	acc Thr	cag Gln 70	gag Glu	ttc Phe	tgg Trp	gat Asp	aac Asn 75	ctg Leu	gaa Glu	aag Lys	gag Glu	aca Thr 80	300
gag Glu	ggc Gly	ctg Leu	agg Arg	cag Gln 85	gag Glu	atg Met	agc Ser	aag Lys	gat Asp 90	ctg Leu	gag Glu	gag Glu	gtg Val	aag Lys 95	gcc Ala	348
aag Lys	gtg Val	cag Gln	ccc Pro 100	tac Tyr	ctg Leu	gac Asp	gac Asp	ttc Phe 105	cag Gln	aag Lys	aag Lys	tgg Trp	cag Gln 110	gag Glu	gag Glu	396
atg Met	gag Glu	ctc Leu 115	tac Tyr	cgc Arg	cag Gln	aag Lys	gtg Val 120	gag Glu	ccg Pro	ctg Leu	cgc Arg	gca Ala 125	gag Glu	ctc Leu	caa Gln	444
gag Glu	ggc Gly 130	gcg Ala	cgc Arg	cag Gln	aag Lys	ctg Leu 135	cac His	gag Glu	ctg Leu	caa Gln	gag Glu 140	aag Lys	ctg Leu	agc Ser	cca Pro	492
ctg Leu 145	ggc Gly	gag Glu	gag Glu	atg Met	cgc Arg 150	gac Asp	cgc Arg	gcg Ala	cgc Arg	gcc Ala 155	cat His	gtg Val	gac Asp	gcg Ala	ctg Leu 160	540
cgc Arg	acg Thr	cat His	ctg Leu	gcc Ala 165	ccc Pro	tac Tyr	agc Ser	gac Asp	gag Glu 170	ctg Leu	cgc Arg	cag Gln	cgc Arg	ttg Leu 175	gcc Ala	588
gcg Ala	cgc Arg	ctt Leu	gag Glu 180	gct Ala	ctc Leu	aag Lys	gag Glu	aac Asn 185	ggc Gly	ggc Gly	gcc Ala	aga Arg	ctg Leu 190	gcc Ala	gag Glu	636
tac Tyr	cac His	gcc Ala 195	aag Lys	gcc Ala	acc Thr	gag Glu	cat His 200	ctg Leu	agc Ser	acg Thr	ctc Leu	agc Ser 205	gag Glu	aag Lys	gcc Ala	684
aag Lys	ccc Pro 210	gcg Ala	ctc Leu	gag Glu	gac Asp	ctc Leu 215	cgc Arg	caa Gln	ggc Gly	ctg Leu	ctg Leu 220	ccc Pro	gtg Val	ctg Leu	gag Glu	732
agc Ser 225	ttc Phe	aag Lys	gtc Val	Ser	ttc Phe 230	ctg Leu	agc Ser	gct Ala	Leu	gag Glu 235	gag Glu	tac Tyr	act Thr	aag Lys	aag Lys 240	780
ctc	aac	acc	cag	tga	ggcg	cccg	cc g	ccgc	cccc	c tt	cccg	gtgc	tca	gaat	aaa	835

Leu Asn Thr Gln * 245

cgtttccaaa gtgggaaaaa aaaa

	<21 <21	.0> 1 .1> 2 .2> E	2454	sapi	lens											
		21> 0	CDS (154)	(1	L710)											
gtga		00> 1 gag g		cgcag	gg tg	gagaa	aagco	g aga	acago	ıcag	gtag	ggaa	at o	gtga	aggtga	•
gtgaaaggag ggaacgcagg tgagaaagcg agacaggcag gtagggaaat cgtgaggtga gcgtgatcct agctccttgt ggcagagcct agagagaagg cgaggacgct gaagaaccag															12	
gcgg	gacag	get <u>e</u>	ggcag	gagag	ga ga	aagtt	egget	ago		: Glı				ı Glı	g cct ı Pro	17
gga Gly	gca Ala	tcc Ser 10	atg Met	gat Asp	gag Glu	aac Asn	tac Tyr 15	ttt Phe	gtg Val	aac Asn	tac Tyr	act Thr 20	ttc Phe	aaa Lys	gat Asp	22
cgg Arg	tca Ser 25	cat His	tca Ser	ggc Gly	cgt Arg	gtg Val 30	gct Ala	caa Gln	ggc Gly	atc Ile	atg Met 35	aaa Lys	ctg Leu	tgt Cys	cta Leu	27
gag Glu 40	gag Glu	gag Glu	ctc Leu	ttt Phe	gct Ala 45	gat Asp	gtc Val	acc Thr	att Ile	tcg Ser 50	gtg Val	gaa Glu	ggc Gly	cgg Arg	gag Glu 55	31
ttt Phe	cag Gln	ctc Leu	cat His	cgg Arg 60	ctg Leu	gtc Val	ctc Leu	tca Ser	gct Ala 65	cag Gln	agc Ser	tgc Cys	ttc Phe	ttc Phe 70	cga Arg	36
tcc Ser	atg Met	ttc Phe	act Thr 75	tcc Ser	aac Asn	ctg Leu	aag Lys	gag Glu 80	gcc Ala	cac His	aac Asn	cgg Arg	gtg Val 85	att Ile	gtg Val	4:
ctg Leu	cag Gln	gat Asp 90	gtc Val	agc Ser	gag Glu	tct Ser	gtt Val 95	ttc Phe	cag Gln	ctc Leu	ctg Leu	gtt Val 100	gat Asp	tat Tyr	atc Ile	4 (
															tat Tyr	51
gag Glu 120	gtg Val	tca Ser	gac Asp	atg Met	tat Tyr 125	cag Gln	ctg Leu	aca Thr	tct Ser	ctc Leu 130	ttt Phe	gag Glu	gaa Glu	tgc Cys	tct Ser 135	5:

	ttt Phe														606
	ctg Leu	_	_		_	-					_	_	_	_	654
	tgt Cys	_	_		_	_	_	_	_				_		702
	cac His 185	_		_				_			_	_		_	750
	tgt Cys														798
	aaa Lys				_		_						_	_	846
_	gaa Glu								_						894
	cgt Arg							_		_	_	_	_	-	942
	atc Ile 265													-	990
	tgc Cys														1038
	cgc Arg														1086
	cct Pro														1134
	aaa Lys														1182
	tcc Ser 345														1230

gag aca act cag cta gag gtg gct gtg tca ggg gct gct ggt gcc aac Glu Thr Thr Gln Leu Glu Val Ala Val Ser Gly Ala Ala Gly Ala Asn 360 365 370 375	278
ctc aac ggg atc atc tac tta cta ggg ggg gag gag aat gat ctg gac 1: Leu Asn Gly Ile Ile Tyr Leu Leu Gly Gly Glu Glu Asn Asp Leu Asp 380 385 390	326
ttc ttt acc aaa cct tcc cga ctc atc cag tgc ttt gac aca gag aca 1 Phe Phe Thr Lys Pro Ser Arg Leu Ile Gln Cys Phe Asp Thr Glu Thr 395 400 405	374
gac aaa tgc cat gtg aag ccc tat gtg ctg ccc ttt gca ggc cgc atg Asp Lys Cys His Val Lys Pro Tyr Val Leu Pro Phe Ala Gly Arg Met 410 415 420	422
cac gca gct gtg cat aaa gat ctg gtg ttc atc gtg gct gaa ggg gac 1 His Ala Ala Val His Lys Asp Leu Val Phe Ile Val Ala Glu Gly Asp 425 430 435	1470
tcc ctg gtg tgc tac aat ccc ttg cta gac agc ttc acc cgg ctt tgc Ser Leu Val Cys Tyr Asn Pro Leu Leu Asp Ser Phe Thr Arg Leu Cys 440 450 455	1518
ctt cct gag gcc tgg agc tct gcc cca tcc ctc tgg aag att gcc agc Leu Pro Glu Ala Trp Ser Ser Ala Pro Ser Leu Trp Lys Ile Ala Ser 460 465 470	1566
tgt aac ggg agc atc tat gtc ttc cgg gac cga tat ada adg ggg 500 Cys Asn Gly Ser Ile Tyr Val Phe Arg Asp Arg Tyr Lys Lys Gly Asp 485	1614
gcc aac acc tac aag ctt gac cct gcc act tca gcc gta act gtc aca Ala Asn Thr Tyr Lys Leu Asp Pro Ala Thr Ser Ala Val Thr Val Thr 490 495 500	1662
aga ggt att aag gtg ctg ctt acc aat ttg cag ttt gtg ttg gcc taa Arg Gly Ile Lys Val Leu Leu Thr Asn Leu Gln Phe Val Leu Ala * 505 510	1710
ggctgtgggg aggggaggag aactgctcac tccttttccc tccccataca aactcaaagt	1770
cccctgggcc ccaattcaga gttatgtttt ttttggcaca tactagaaag gcagtgcctc	1830
agcccttccc tgaatccatg gaggtgttct gtttggggct ttttagactg ctgctgctca	1890
gctggttgct tgaactgaca gtaggccagc ctgttctctg ccattcccta gtcatcctgt	1950
gcctcaccac agcttgctta gagcaagcct tttctcagac cttaggcaca gcctctcctc	2010
tttacctgat caatgttaaa tgtaagcacc cctgatccca ggacataagg aaagatgccc	2070
aattgtactt ttgttctata gcctgtgaaa tggctagttg atcatttttc cacaaagaat	2130
taggtgttaa gagttttcct tcaggcttta cttaggagaa tggactaagc tgaaggtgta	2190
cttcaccagc aagagtcaac tctagaattc aggatgttcc ttctattgtt ttcttagcca	2250

tctgtcagga	aatgtaactt	tggttttatt	tttggcttat	tccaaggggt	aagccagaaa	2310
atagaaatga	ttatttctga	ttaatagcag	aaacttttc	aatctcaaat	atataaggtg	2370
tctgctcttt	taaaagctct	aagctaagtc	aagagctagg	aactgttgat	acaaataaaa	2430
gtttttgaag	ggaaaaaaaa	aaaa				2454

<210> 19 <211> 3586 <212> DNA <213> Homo sapiens													
<220> <221> CDS <222> (106)(1860)													
<400> 19 gtaccggtcc ggaattcccg ggtcgacgat ttcgtggagc aggcagacgt cgaagccgtc	60												
ctgcagccga ccctcctgca tctgggcacg ttcaggttgc gacac atg aag gct Met Lys Ala 1													
ttg atg aac gag aag gcc cag gcc gcc ctg gtg gag ttt gtg gag gac Leu Met Asn Glu Lys Ala Gln Ala Ala Leu Val Glu Phe Val Glu Asp 5 10 15	162												
gtc aat cac gct gcc att ccc agg gag atc cca cgc aag gat ggg gtc Val Asn His Ala Ala Ile Pro Arg Glu Ile Pro Arg Lys Asp Gly Val 20 25 30 35	210												
tgg agg gtt ctg tgg aag gac cgt gcg cag gac acg agg gtc ctg agg Trp Arg Val Leu Trp Lys Asp Arg Ala Gln Asp Thr Arg Val Leu Arg 40 45 50	258												
cag atg aca cgc ctg ctg ctg gat gac ggg ccc acg cag gcc gcg gag Gln Met Thr Arg Leu Leu Leu Asp Asp Gly Pro Thr Gln Ala Ala Glu 55 60 65	306												
gct ggg acc ccc ggt gag gca ccc acc cct ccc gct tcg gag acg cag Ala Gly Thr Pro Gly Glu Ala Pro Thr Pro Pro Ala Ser Glu Thr Gln 70 75 80	354												
gcc cag gat tct ggt gag gta aca ggg cat gct ggc tcg ctt ctt ggg Ala Gln Asp Ser Gly Glu Val Thr Gly His Ala Gly Ser Leu Leu Gly 85 90 95	402												
gca ccc agg aac cca agg agg ggc cgt cgg ggt cgc aga aac aga acc Ala Pro Arg Asn Pro Arg Arg Gly Arg Arg Gly Arg Arg Asn Arg Thr 100 105 110 115	450												
aga cgc aac agg ttg acc cag aag ggc aag aag aga agc cga gga g	498												

·	
Arg Arg Asn Arg Leu Thr Gln Lys Gly Lys Lys Arg Ser Arg Gly Gly 120 125 130	
cgg ccg tct gct ccc gcg agg agt gag gcc gag gac tct tcc gac gag Arg Pro Ser Ala Pro Ala Arg Ser Glu Ala Glu Asp Ser Ser Asp Glu 135	546
agc ctg ggc atc gtg atc gag gag atc gac cag ggc gac ctg agc gga Ser Leu Gly Ile Val Ile Glu Glu Ile Asp Gln Gly Asp Leu Ser Gly 150 155 160	594
gaa gag gac cag agc gcg ctg tac gcc acg ctg cag gcc gct gcc agg Glu Glu Asp Gln Ser Ala Leu Tyr Ala Thr Leu Gln Ala Ala Ala Arg 165 170 175	642
gag ctg gtt agg cag tgg gcg ccc tgc aac tcc gag ggg gaa gaa gac Glu Leu Val Arg Gln Trp Ala Pro Cys Asn Ser Glu Gly Glu Glu Asp 180 185	690
ggt ccc cgc gag ttc ttg gct ctg gtc acc gtc acc gac aaa tcg aag Gly Pro Arg Glu Phe Leu Ala Leu Val Thr Val Thr Asp Lys Ser Lys 200 205	738
aaa gaa gag gca gag aag gag cca gct ggg gcc gaa tcc atc cgc ttg Lys Glu Glu Ala Glu Lys Glu Pro Ala Gly Ala Glu Ser Ile Arg Leu 225	786
aac acc aaa gaa gac aaa aat ggt gtc ccc gac tta gtg gcc ctg ctg Asn Thr Lys Glu Asp Lys Asn Gly Val Pro Asp Leu Val Ala Leu Leu 230 235	834
gct gtg aga gac acc ccg gac gag gag ccg gtg gac agc gac act tcg Ala Val Arg Asp Thr Pro Asp Glu Glu Pro Val Asp Ser Asp Thr Ser 245 250 255	882
gag agc gac tcg cag gaa agt ggg gac caa gaa aca gag gag ttg gat Glu Ser Asp Ser Gln Glu Ser Gly Asp Gln Glu Thr Glu Glu Leu Asp 275 260 275	930
aat cct gag ttc gtg gcc att gtg gcc tat acc gac ccg tcg gac ccc Asn Pro Glu Phe Val Ala Ile Val Ala Tyr Thr Asp Pro Ser Asp Pro 280 285 290	978
tgg gcc cgg gag gag atg ttg aaa atc gct tct gtt atc gag tcg ctg Trp Ala Arg Glu Glu Met Leu Lys Ile Ala Ser Val Ile Glu Ser Leu 305	1026
ggc tgg agc gac gag aaa gac aag cga gac ccc ctc cga cag gtc ttg Gly Trp Ser Asp Glu Lys Asp Lys Arg Asp Pro Leu Arg Gln Val Leu 310	1074
tcc gtc atg tcc aag gac act aac ggg acc cgc gtg aag gtg gaa gag Ser Val Met Ser Lys Asp Thr Asn Gly Thr Arg Val Lys Val Glu Glu 325 330	1122
gcg ggc cgc gag gtg gac gcc gtg gtc ctg cgc aag gcc ggg gat gac Ala Gly Arg Glu Val Asp Ala Val Val Leu Arg Lys Ala Gly Asp Asp	1170

340	345	350	355
ggg gac ctc cgg gag Gly Asp Leu Arg Glo 360	ı Cys Ile Ser Thr	ttg gcg cag ccg gat Leu Ala Gln Pro Asp 365	ctc cct 1218 Leu Pro 370
ccc cag gcg aag aag Pro Gln Ala Lys Lys 375	g gct ggg cgt ggc s Ala Gly Arg Gly 380	ctc ttc ggg ggc tgg Leu Phe Gly Gly Trp 385	agc gag 1266 Ser Glu
cac cgt gag gac gad His Arg Glu Asp Glu 390	a ggg ggt ctt ctg u Gly Gly Leu Leu 395	gag ctg gtg gcg ctc Glu Leu Val Ala Leu 400	ctg gct 1314 Leu Ala
gcc cag gac atg gc Ala Gln Asp Met Al 405	g gag gtg atg aag a Glu Val Met Lys 410	gag gaa aaa gaa aac Glu Glu Lys Glu Asn 415	gcc tgg 1362 Ala Trp
gaa ggc ggg aag ta Glu Gly Gly Lys Ty 420	c aaa tac ccc aaa r Lys Tyr Pro Lys 425	ggc aaa ctg ggg gag Gly Lys Leu Gly Glu 430	gta ttg 1410 Val Leu 435
gcg ctc ctg gcc gc Ala Leu Leu Ala Al 44	a Arg Glu Asn Met	ggg tcc aac gag ggg Gly Ser Asn Glu Gly 445	tcg gag 1458 Ser Glu 450
gag gct tcg gac ga Glu Ala Ser Asp Gl 455	a cag tcc gag gag u Gln Ser Glu Glu 460	gag tcg gag gac acc Glu Ser Glu Asp Thr 465	gag agc 1506 Glu Ser
gag gcg tcg gag cc Glu Ala Ser Glu Pr 470	g gag gac agg gca o Glu Asp Arg Ala 475	tcc agg aag ccc cgg Ser Arg Lys Pro Arg 480	gcc aag 1554 Ala Lys
agg gcg cgc acg gc Arg Ala Arg Thr Al 485	c ccc agg ggc ctg a Pro Arg Gly Leu 490	act ccg gcc ggc gcg Thr Pro Ala Gly Ala 495	cct ccc 1602 Pro Pro
		gcg ggc ggc cga ggc Ala Gly Gly Arg Gly 510	
cga ggc cgg ggc gt Arg Gly Arg Gly Va 52	l Thr Pro Glu Lys	aaa gcc ggg agc cgg Lys Ala Gly Ser Arg 525	ggc tcg 1698 Gly Ser 530
gcc cag gac gac gc Ala Gln Asp Asp Al 535	c gca gga agc agg a Ala Gly Ser Arg 540	aag aag aag ggg agc Lys Lys Lys Gly Ser 545	gcg ggc 1746 Ala Gly
tcc ggg gcc cat gc Ser Gly Ala His Al 550	c agg gca ggc gag a Arg Ala Gly Glu 555	gcc aag ggc cag gcg Ala Lys Gly Gln Ala 560	ccc act 1794 Pro Thr
		gcc cgt cgg ggc cgg Ala Arg Arg Gly Arg 575	

ccc cct aaa tgc cgc tag tggccc cccaagaagc cgcccaggct gcgagcaggc Pro Pro Lys Cys Arg * 580 585	1896
cccgcagggc acccgcccgc ctgtggcccc cgccctcccc tcccctcttc ctgtcctccg	1956
cagacgcaat ctcctcgctt cacagcgcgc ccgggccgcg ttttgccagc gtcacgttcc	2016
cctctcgggc cctcgcaggc cgggggcgcc agcgatcccg acggaggaag cccggatggg	2076
aggaggaaag agaagtgggc gcccgaggca gcagcgcagg gccgagatgg ggacgcgcca	2136
agtggaccag gattgggggc ccgggttgcc cccggagggg gtgtgtgtgt ggacgccggg	2196
cacetgeaga ggegageagg getettegtg gegetetegg ggeetgegee tggeaggtge	2256
tgtaggccgc tgtcgcccct accccagtct gactgggccc tgggtctgtg gtggaggctc	2316
agtcaccagc cgcgcagcgc gtgtcagggc gcaactctca gccaggggag gccccagctc	2376
ccagccaggg gaagagatga ttccagaaag gaaagtctga gagatagaag gcggttggga	2436
aggggaggag gaggaaaggg gagaggaacg gtgggagaag ggaaagagga ggaggagggg	2496
gaggggggag cagagggaag acacatgcca gccctgccta ctggggcgcc cctgataaca	2556
aaggaaccag ccccaggcca aggccacctg cccctgacca caagttgaat ttgtcactca	2616
gactgcagtg tttcccaaca ttctaattat ttgcagaggt gttcaatttg gggtaattca	2676
cttaaaatcc agttttggtt cttctgggct gagtgggccc tggcccctcc cataggctgt	2736
ggctcccctg ggtgccccct ctccagtgga gctgacccac cgctcagcgc tggccttgca	2796
gcccttacta aaagacttga aagtccctgg gttcaccccc tgagtgaatt aaaggccaga	2856
ggggccccga agggcactgt gagggacaga ggctcacctg ggcagtgcag aagccggccg	2916
cgtgtccctc cttacagggg atgaaatgac ctggggagga aaccccagcc ctgccctgga	2976
ggttccagag taggcgggcc ggtgctgtga ggcttcacaa cctgctgtcc caagcacgct	3036
tgagttgtat gtgagtetgt geegtgeegt geegtatget teageteetg eaaceeegge	3096
tgagetegat ttttaeetaa atateagtet eeaegggaee eeaeetteat teatgeette	3156
ttgtccctgg ggcaatgtgt gtgcttcctc gtcccaattt ccattccctg gcagtgagga	3216
gcccatcgtg ccagggggcc ctgccccact tgtccctggg aaggaatagg agggtttggg	3276
tgtgacctca cagtccagac cagactgtcc cagtcctatg tcagggacac ccagatgtag	3336
aagctgactg agacctgctg cagggcgtgg gtgctcccct ctgcttggag gctgtccctg	3396
gacagtgacc cacccactga ggaccaggct gggtgtacct tgagctgggc acagcagcct	3456

	gtggt	tgtt	gc ct	tgtg	ggtg	g gg:	aggg	cccc	agg	tgtg	ctt	ctcc	gtag	gc a	gtcc	taggc	3516
	ttct	ctcc	ct g	tgcc	ctgt	g tc	acct	ggat	cct	ccag	taa	agtga	aaati	tc a	gcac	tgtaa	3576
	aaaa	aaaaaaaaaa														3586	
			•														
			0 > 2 1 > 3														
			2> D 3> H		sapi	ens											
		<22	0 >														
			1> C 2> (. (18	94)											
		<pre><400> 20 gatacagttc tgaccatcat tacgccaatc ttggcacgag gggaag atg gcg gag</pre>													==		
	gata	cagt	tc t	gacc	atca	t ta.	cgcc	aato	ttg	gcac	gag	ggga	ag		Ala	gag Glu	55
Son that you bent bodd then the side that the sont the sont that the sont that the sont the sont the sont that the sont the so	tcc Ser	ggc Gly 5	ggt Gly	agc Ser	agc Ser	ggt Gly	ggt Gly 10	gct Ala	ggt Gly	ggc Gly	ggc Gly	ggc Gly 15	gct Ala	ttc Phe	ggc Gly	gcg Ala	103
11	ggc Gly 20	ccg Pro	ggc Gly	ccc Pro	gag Glu	cgc Arg 25	ccg Pro	aac Asn	agc Ser	acg Thr	gcc Ala 30	gac Asp	aag Lys	aac Asn	ggg ggg	gcc Ala 35	151
reg H Hung H H	ctc Leu	aag Lys	tgc Cys	acc Thr	ttc Phe 40	tcg Ser	gca Ala	ccc Pro	agc Ser	cac His 45	agc Ser	acc Thr	agc Ser	ctc Leu	ctg Leu 50	cag Gln	199
	ggc Gly	ctg Leu	gcc Ala	acc Thr 55	ctc Leu	cgc Arg	gct Ala	cag Gln	ggc Gly 60	cag Gln	ctc Leu	ctc Leu	gat Asp	gtt Val 65	gtg Val	ctg Leu	247
	act Thr	att Ile	aac Asn 70	aga Arg	gag Glu	gcc Ala	ttt Phe	cct Pro 75	gca Ala	cac His	aag Lys	gtc Val	gtc Val 80	ctg Leu	gct Ala	gcc Ala	295
	tgc Cys	agc Ser 85	Asp	tac Tyr	ttc Phe	agg Arg	gcc Ala 90	atg Met	ttc Phe	acc Thr	ggc Gly	ggc Gly 95	atg Met	cgg Arg	gag Glu	gca Ala	343
	agc Ser 100	Gln	gac Asp	gtc Val	atc Ile	gag Glu 105	Leu	aag Lys	ggc Gly	gtg Val	tcg Ser 110	gcc Ala	cgt Arg	ggc Gly	ctg Leu	cgg Arg 115	391
	cac His	atc Ile	atc Ile	gac Asp	ttc Phe 120	Ala	tac Tyr	agc Ser	gcc Ala	gag Glu 125	gtg Val	aca Thr	ctg Leu	gac Asp	ctg Leu 130	Asp	439
	tgc Cys	gtg Val	cag Gln	gac Asp	gtg Val	ctg Leu	ggc Gly	gcg	gcc Ala	gtg Val	ttc Phe	ttg Leu	cag Gln	atg Met	ctg Leu	ccc	487

135 140 145 535 gtg gtg gag ctg tgc gag gag ttc ctg aag gcg gcc atg agc gtg gag Val Val Glu Leu Cys Glu Glu Phe Leu Lys Ala Ala Met Ser Val Glu 150 155 583 acc tgc ctc aac atc ggc cag atg gcc acc acc ttc agc ctg gcc tcg Thr Cys Leu Asn Ile Gly Gln Met Ala Thr Thr Phe Ser Leu Ala Ser 165 170 175 ctg cga gag tcg gtg gat gcc ttc acc ttc cgg cac ttc ctg caq atc 631 Leu Arg Glu Ser Val Asp Ala Phe Thr Phe Arg His Phe Leu Gln Ile 180 185 190 gcc gag gag gat ttc ctg cgc ctg cca ctg gag cgc ctg gtc ttc 679 Ala Glu Glu Glu Asp Phe Leu Arg Leu Pro Leu Glu Arg Leu Val Phe 200 210 ttc ctg cag agc aac cgg ctg cag agc tgt gcc gag atc gac ctg ttc 727 Phe Leu Gln Ser Asn Arg Leu Gln Ser Cys Ala Glu Ile Asp Leu Phe 775 ege geg gee gte ege tgg etg eag eat gae eeg gee egg egg eeg ege Arg Ala Ala Val Arg Trp Leu Gln His Asp Pro Ala Arg Arg Pro Arg gee age cae gtg etc tge cae att ege tte eeg etc atg eag teg tee 823 Ala Ser His Val Leu Cys His Ile Arg Phe Pro Leu Met Gln Ser Ser 250 gag ctg gtg gac agc gtg cag acg ctg gac atc atg gtg gag gac gtg 871 Glu Leu Val Asp Ser Val Gln Thr Leu Asp Ile Met Val Glu Asp Val 260 265 270 ctg tgc cgc cag tat ctg ctg gag gcc ttc aac tac cag gtg ctg ccc 919 Leu Cys Arg Gln Tyr Leu Leu Glu Ala Phe Asn Tyr Gln Val Leu Pro 280 ttc cgg cag cac gag atg cag tct ccg cgc acc gcc gtg cgc tcg gat 967 Phe Arg Gln His Glu Met Gln Ser Pro Arg Thr Ala Val Arg Ser Asp 295 gtg ccc tcg ctc gtc acc ttc ggc ggc acg ccc tac acc gac agc gac 1015 Val Pro Ser Leu Val Thr Phe Gly Gly Thr Pro Tyr Thr Asp Ser Asp 310 1063 cgc tcg gtc agc agc aag gtc tac cag ctg cct gag ccg gga gcc cgc Arg Ser Val Ser Ser Lys Val Tyr Gln Leu Pro Glu Pro Gly Ala Arg cac ttc cgc gag ctc acg gag atg gag gta ggc tgc agc cac acg tgc 1111 His Phe Arg Glu Leu Thr Glu Met Glu Val Gly Cys Ser His Thr Cys 345 gtg gcc gtg ctg gac aat ttt gtg tac gtg gcc ggg ggg cag cac ctg 1159 Val Ala Val Leu Asp Asn Phe Val Tyr Val Ala Gly Gly Gln His Leu 360

cag Gln	tac Tyr	cgc Arg	agc Ser 375	ggc Gly	gag Glu	ggc Gly	gca Ala	gtg Val 380	gac Asp	gcc Ala	tgc Cys	tac Tyr	cgc Arg 385	tac Tyr	gac Asp	120)7
ccc Pro	cac His	ctg Leu 390	aat Asn	cgc Arg	tgg Trp	ctg Leu	cgc Arg 395	ctg Leu	cag Gln	gcc Ala	atg Met	cag Gln 400	gaa Glu	agc Ser	cgc Arg	125	55
atc Ile	cag Gln 405	ttc Phe	cag Gln	ctg Leu	aac Asn	gtg Val 410	ctg Leu	tgc Cys	ggc Gly	atg Met	gtg Val 415	tac Tyr	gcc Ala	acg Thr	ggc Gly	130)3
ggc Gly 420	cgc Arg	aac Asn	cga Arg	gcc Ala	ggc Gly 425	agc Ser	ctg Leu	gcc Ala	tcc Ser	gtg Val 430	gag Glu	cgg Arg	tac Tyr	tgc Cys	ccc Pro 435	135	51
cgg Arg	cgc Arg	aat Asn	gag Glu	tgg Trp 440	ggc Gly	tac Tyr	gcc Ala	tgc Cys	tcg Ser 445	ctg Leu	aag Lys	cgc Arg	cgt Arg	acc Thr 450	tgg Trp	139	∍9
ggc Gly	cat His	gct Ala	999 Gly 455	gcc Ala	gcc Ala	tca Ser	gly aaa	ggc Gly 460	cgc Arg	ctc Leu	tac Tyr	atc Ile	tcg Ser 465	ggt Gly	ggc Gly	144	47
tac Tyr	gly 999	atc Ile 470	tca Ser	gtg Val	gag Glu	gac Asp	aag Lys 475	aag Lys	gcc Ala	ctg Leu	cac His	tgc Cys 480	tac Tyr	gac Asp	ccc Pro	14:	95
gtg Val	gcc Ala 485	gac Asp	cag Gln	tgg Trp	gag Glu	ttc Phe 490	aag Lys	gcg Ala	ccc Pro	atg Met	agc Ser 495	gaa Glu	ccc Pro	cgc Arg	gtg Val	15	43
cta Leu 500	cac His	gcc Ala	atg Met	gtg Val	ggt Gly 505	gcc Ala	ggc Gly	ggc Gly	cgc Arg	atc Ile 510	tat Tyr	gcc Ala	ctc Leu	ggg ggg	ggc Gly 515	15	91
cgc Arg	atg Met	gac Asp	cac His	gcg Ala 520	gac Asp	cgc Arg	tgc Cys	ttc Phe	gac Asp 525	gtg Val	ctg Leu	gct Ala	gtg Val	gag Glu 530	tac Tyr	16	39
tat Tyr	gtg Val	ccg Pro	gag Glu 535	Thr	gac Asp	cag Gln	tgg Trp	acc Thr 540	agc Ser	gtg Val	agc Ser	ccc Pro	atg Met 545	cgg Arg	gcc Ala	16	87
ggc Gly	cag Gln	tca Ser 550	Glu	gcc Ala	ggc Gly	tgc Cys	tgc Cys 555	ctg Leu	ctg Leu	gag Glu	agg Arg	aag Lys 560	atc Ile	tac Tyr	atc Ile	17	35
gtc Val	999 Gly 565	Gly	tac Tyr	aac Asn	tgg Trp	cgt Arg 570	Leu	aac Asn	aac Asn	gtc Val	acg Thr 575	ggc Gly	atc Ile	gta Val	cag Gln	17	83
gtg Val 580	Tyr	aac Asn	acg Thr	gac Asp	acc Thr 585	Asp	gag Glu	tgg Trp	gag Glu	cgg Arg 590	Asp	ctg Leu	cac His	ttc Phe	ccg Pro 595	18	31

gag tcc tt Glu Ser Ph	c gca ggc ne Ala Gly 600	ata gcc Ile Ala	tgc gcc Cys Ala	e ccc a Pro 605	gtc (Val 1	ctg ctg Leu Leu	ccc Pro	cgg Arg 610	gcc Ala	1879
ggg acc ag Gly Thr Ar	gg agg tag gg Arg * 615	ccccaag	gac ccc	egggac	c ctg	ggcctga	c cg	catgt	tgt	1934
ctccaagtgg	ggettgge	ga atgcad	gtct go	cctgaga	aac o	ccagtg	ccc (ccctt	cgccc	1994
gggctgccct	tgaggggc	ct gctgcg	gttga ta	agccc	ccc t	cccagg	ggt (ccctc	cctcc	2054
ctccttccca	aagcagato	cc tggctg	gcgag to	catcc	gag g	ggagcct	gcc (ggcaa	agcgt	2114
ctgacatgtg	gtggcagca	a attcgt	cccc gg	ggtggt	tt c	ectegee	tgg (cccc	gagtc	2174
cccacgggct	ggcgggtgg	ga atccca	iggto to	cagggg	ggt c	cctgtg	cag o	ctcca	tctca	2234
cttctctact	gcctcccag	c cccacg	ıgttt ca	ıggcatt	ca g	gatgtgag	gct d	catca	acatt	2294
gaacccaaag	teggtggta	ıt atgact	cacc ct	ccttcc	aa g	gtctcctg	gcg (cgcgt	gtttt	2354
taaaataaac	tcacccgaa	a cgtccg	rtaac ac	acggad	ct c	caggago	cag t	gaga	ggtgg	2414
cttagaagcc	cctggtttg	g ggtggg	tgga gg	aggagg	ıgc a	ıcgtgtct	gc d	ctccc	ctggg	2474
gtgccctcct	tcccccatc	c caagct	gctg ag	gggagg	icc c	tggtcat	gc c	ctcag	ttcct	2534
gccttcatct	gctttccgg	a ggaaaa	acca ta	tcaact	cc t	agaaacg	get c	ctta	ggggc	2594
ttgggacctt	ccatttggc	a ctgagc	atct tg	tggggc	ct t	aactggo	ctg a	igaca	tcccg	2654
gccctctaca	tttgccctg	t tggcca	ggca gt	cccctt	.cc c	gcaatto	gga g	ggcg	acgct	2714
aacttcagaa	tcccatagt	g gcgctt	gccg ca	ggtctg	gt g	gggtgto	tt t	tttc	tcctc	2774
cctcctttcc	acccctccg	c gccctg	ccac tc	cctggc	ct g	ccctgtt	tt t	gggt	caaca	2834
ttgctacgga	gccagcagt	g aggcct	ttcc ct	tcaagg	gc t	ctgtggt	at c	tctg	gccac	2894
atttgttcta	atgtctgaa	c ctctaa	atct tt	tctttt	tg a	ttggttt	ta c	tgtt	tttaa	2954
gaagccagca	ctgctgtct	c atagat	ggga tt	tgtact	ct t	ggggcaa	ct t	aaagt	gtct	3014
ctctcgctgc	taacagacg	a ttgatg	tctt gt	ctctgt	ga c	ccactca	cc a	tgtaa	aagaa	3074
ttaacctcct	atcttagca	g acatcg	tctc ct	aatatt	tc c	ctttatt	ta a	taaaa	aatgt	3134
tatggtgaag	agatggagc	c ggccca	gcac tg	agcttg	tg c	ggcttgg	gt c	tgatt	ggtc	3194
acagattcct	cgtgtgtcc	t ccgcgt	gtct gg	gggctc	ct c	tcccccg	cc t	cagco	ctccc	3254
cccgcatccc	cccaagaaa	g gaaaat	tatt tt	tcgtat	tg ta	aaacttt	aa a	catga	aaaa	3314
gctgtttta	atttaaaaa	a aaaaaa				·				3340

<210> 21 <211> 2547 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (143)..(2275) <400> 21 aaggateett aattaaatta ateeeeeee eeeeeegee eeegeggtgg eggeggegge 60 ggcggtggct gccgtggcgg ctgagagtcc agagccggac gttccggccg cttcgggctg 120 geggetggag agegeteggg to atg tet gee cag ggg gae tge gag tte etg 172 Met Ser Ala Gln Gly Asp Cys Glu Phe Leu gtg cag cga gcc cgg gag ttg gtg ccg caa gac ctg tgg gca gcc aag 220 Val Gln Arg Ala Arg Glu Leu Val Pro Gln Asp Leu Trp Ala Ala Lys geg tgg etg ate acg gec ege age etc tac eeg gea gae ttt aac ate 268 Ala Trp Leu Ile Thr Ala Arg Ser Leu Tyr Pro Ala Asp Phe Asn Ile 30 cag tat gag atg tac acc atc gag cgg aat gca gag cgg acc gcc acc 316 Gln Tyr Glu Met Tyr Thr Ile Glu Arg Asn Ala Glu Arg Thr Ala Thr 50 45 gcc ggg agg ctg ctg tac gac atg ttt gtg aat ttc cca gac cag ccg 364 Ala Gly Arg Leu Leu Tyr Asp Met Phe Val Asn Phe Pro Asp Gln Pro 65 60 gtg gtg tgg aga gaa atc agc att att aca tca gca tta agg aac gat 412 Val Val Trp Arg Glu Ile Ser Ile Ile Thr Ser Ala Leu Arg Asn Asp 85 80 75 tca cag gac aaa caa acc caa ttt tta aga agt tta ttt gaa act ctt 460 Ser Gln Asp Lys Gln Thr Gln Phe Leu Arg Ser Leu Phe Glu Thr Leu 100 95 cct ggt cgg gtc cag tgt gaa atg tta cta aag gtc acg gaa caa tgc 508 Pro Gly Arg Val Gln Cys Glu Met Leu Leu Lys Val Thr Glu Gln Cys 120 115 110 ttc aac acg tta gaa cga tca gaa atg ttg ctt cta ctt ttg agg cgc 556 Phe Asn Thr Leu Glu Arg Ser Glu Met Leu Leu Leu Leu Arg Arg 130 125 ttc cct gaa acg gtg gtg cag cat ggg gtt ggc ctt ggg gag gca cta 604 Phe Pro Glu Thr Val Val Gln His Gly Val Gly Leu Gly Glu Ala Leu 150 652 tta gag gct gaa act att gaa gaa caa gaa tct cca gtg aac tgc ttt

Leu 155		ı Ala	Glu	Thr	Ile		Glu	Gln	Glu	Ser 165		Val	Asn	Cys	Phe 170	
					Cys							att :Ile				700
gat Asp	gtt Val	cga Arg	tta Leu 190	Pro	gcc Ala	aat Asn	tta Leu	ttg Leu 195	Tyr	aag Lys	tac Tyr	ttg Leu	aac Asn 200	Lys	gca Ala	748
gct Ala	gaa Glu	ttt Phe 205	tat Tyr	atc Ile	aat Asn	tat Tyr	gtc Val 210	act Thr	agg Arg	tct Ser	act Thr	caa Gln 215	ata Ile	gaa Glu	aat Asn	796
cag Gln	cat His 220	caa Gln	ggc Gly	gcc Ala	cag Gln	gat Asp 225	aca Thr	tct Ser	gat Asp	tta Leu	atg Met 230	tca Ser	cct Pro	agc Ser	aaa Lys	844
cgt Arg 235	agc Ser	tct Ser	cag Gln	aag Lys	tac Tyr 240	ata Ile	ata Ile	gaa Glu	ggg Gly	ctg Leu 245	acg Thr	gaa Glu	aaa Lys	tca Ser	tcc Ser 250	892
												ttg Leu				940
gga Gly	atg Met	aga Arg	tgt Cys 270	gaa Glu	tgg Trp	cag Gln	atg Met	gat Asp 275	aaa Lys	gga Gly	aga Arg	cga Arg	agc Ser 280	tat Tyr	gga Gly	988
gat Asp	att Ile	ttg Leu 285	cat His	aga Arg	atg Met	aag Lys	gat Asp 290	ctc Leu	tgc Cys	aga Arg	tac Tyr	atg Met 295	aac Asn	aac Asn	ttt Phe	1036
gat Asp	agt Ser 300	gaa Glu	gca Ala	cat His	gca Ala	aaa Lys 305	tat Tyr	aaa Lys	aac Asn	caa Gln	gtg Val 310	gtg Val	tat Tyr	tcc Ser	acc Thr	1084
atg Met 315	ctg Leu	gtc Val	ttc Phe	ttt Phe	aag Lys 320	aat Asn	gca Ala	ttc Phe	cag Gln	tat Tyr 325	gtc Val	aac Asn	agc Ser	ata Ile	cag Gln 330	1132
cca Pro	tct Ser	ctc Leu	ttc Phe	caa Gln 335	ggt Gly	cct Pro	aat Asn	gcc Ala	ccg Pro 340	agc Ser	caa Gln	gtt Val	cca Pro	ctg Leu 345	gtt Val	1180
ctt Leu	ctt Leu	gaa Glu	gat Asp 350	gta Val	tcg Ser	aat Asn	gtg Val	tat Tyr 355	ggt Gly	gat Asp	gta Val	gaa Glu	att Ile 360	gat Asp	cgt Arg	1228
aat Asn	aaa Lys	cac His 365	atc Ile	cat His	aaa Lys	aag Lys	agg Arg 370	aaa Lys	cta Leu	gct Ala	gaa Glu	gga Gly 375	aga Arg	gaa Glu	aaa Lys	1276
acc Thr	atg Met	agt Ser	tca Ser	gac Asp	gat Asp	gaa Glu	gac Asp	tgt Cys	tcg Ser	gcg Ala	aaa Lys	gga Gly	aga Arg	aat Asn	cgt Arg	1324

390 380 cac att gta gtc aat aaa gcc gaa ctt gct aac tcc act gaa gtg tta 1372 His Ile Val Val Asn Lys Ala Glu Leu Ala Asn Ser Thr Glu Val Leu 405 400 395 gaa agc ttt aaa ttg gcc agg gag agc tgg gag ttg ctc tat tcc cta 1420 Glu Ser Phe Lys Leu Ala Arg Glu Ser Trp Glu Leu Leu Tyr Ser Leu gaa ttc ctt gac aaa gaa ttt aca agg att tgc ttg gcc tgg aag acg 1468 Glu Phe Leu Asp Lys Glu Phe Thr Arg Ile Cys Leu Ala Trp Lys Thr 430 gat act tgg ctt tgg tta aga atc ttc ctc act gat atg atc atc tat 1516 Asp Thr Trp Leu Trp Leu Arg Ile Phe Leu Thr Asp Met Ile Ile Tyr 445 cag ggt caa tat aaa aag gcg ata gcc agc ctg cat cac tta gca gct 1564 Gln Gly Gln Tyr Lys Lys Ala Ile Ala Ser Leu His His Leu Ala Ala ctc cag gga tcc att tct cag cca cag atc aca ggg cag ggg acc ctg 1612 Leu Gln Gly Ser Ile Ser Gln Pro Gln Ile Thr Gly Gln Gly Thr Leu gag cat cag agg gcg ctc atc cag ctg gcg acg tgc cac ttt gcg cta 1660 Glu His Gln Arg Ala Leu Ile Gln Leu Ala Thr Cys His Phe Ala Leu 495 ggg gag tac aga atg aca tgt gaa aaa gtc ctt gat ttg atg tgc tac 1708 Gly Glu Tyr Arg Met Thr Cys Glu Lys Val Leu Asp Leu Met Cys Tyr 510 atg gta ctc ccc att caa gat gga ggc aaa tcc cag gag gaa ccc tcg 1756 Met Val Leu Pro Ile Gln Asp Gly Gly Lys Ser Gln Glu Glu Pro Ser 530 525 aaa gta aag ccc aaa ttt aga aaa ggt tcg gat ctg aag ctc ctg cct 1804 Lys Val Lys Pro Lys Phe Arg Lys Gly Ser Asp Leu Lys Leu Pro 545 540 tgt acc agc aag gct atc atg cca tac tgc ctc cat tta atg tta gcc 1852 Cys Thr Ser Lys Ala Ile Met Pro Tyr Cys Leu His Leu Met Leu Ala 565 560 555 tgt ttt aag ctt aga gct ttc aca gac aac aga gac gac atg gca ttg 1900 Cys Phe Lys Leu Arg Ala Phe Thr Asp Asn Arg Asp Asp Met Ala Leu 575 ggg cat gtg att gtg ttg ctt cag caa gag tgg cca cgg ggc gag aat 1948 Gly His Val Ile Val Leu Leu Gln Gln Glu Trp Pro Arg Gly Glu Asn 590 ctt ttc ctg aaa gct gtc aat aaa att tgc caa caa gga aat ttc caa 1996 Leu Phe Leu Lys Ala Val Asn Lys Ile Cys Gln Gln Gly Asn Phe Gln 605

tat gag aat ttt ttc aat tac gtt aca aat att gat atg ctg gag gaa Tyr Glu Asn Phe Phe Asn Tyr Val Thr Asn Ile Asp Met Leu Glu Glu 620 625 630	2044
ttt gcc tac ttg aga act cag gaa ggt ggg aaa att cat ctg gaa tta Phe Ala Tyr Leu Arg Thr Gln Glu Gly Gly Lys Ile His Leu Glu Leu 635 640 645 650	2092
cta ccc aat caa gga atg ctg atc aag cac cac act gta act cga ggc Leu Pro Asn Gln Gly Met Leu Ile Lys His His Thr Val Thr Arg Gly 655 660 665	2140
atc acc aaa ggc gtg aag gag gac ttt cgc ctg gcc atg gag cgc cag Ile Thr Lys Gly Val Lys Glu Asp Phe Arg Leu Ala Met Glu Arg Gln 670 675 680	2188
gtc tcc cgc tgt gga gag aat ctg atg gtg gtt ctg cac agg ttc tgc Val Ser Arg Cys Gly Glu Asn Leu Met Val Val Leu His Arg Phe Cys 685 690 695	2236
att aat gag aag atc ttg ctc ctt cag act ctg acc tga gtggagacct Ile Asn Glu Lys Ile Leu Leu Leu Gln Thr Leu Thr * 700 705 710	2285
ttccaccaga cacagctcgg gcctgtgtaa ttgtaggaga agacactcag cagtgattgc	2345
catggcacag agccgtggtc attgttgctg ttacaaagaa gaaaaccatc tgagttctaa	2405
ctccttggtt gcttaaaagt agttcccaag agtctgagaa gctatttcta tttttaagag	2465
tcattttttg taatttttgt aaaacaaaag taccaatctg ttttgtaaat aaaaatcatc	2525
ctaaaatttg aaaaaaaaa aa	2547

<210> 22

<211> 2625

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (143)..(2353)

<400> 22

aaggatcett aattaaatta atccccccc ccccccgcc cccgcggtgg cggcggcggc 60
ggcggtggct gccgtggcgg ctgagagtcc agagccggac gttccggccg cttcgggctg 120
gcggctggag agcgctcggg tc atg tct gcc cag ggg gac tgc gag ttc ctg
Met Ser Ala Gln Gly Asp Cys Glu Phe Leu
1 5 10

gtg cag cga gcc cgg gag ttg gtg ccg caa gac ctg tgg gca gcc aag

Val	Gln	Arg	Ala	Arg 15	Glu	Leu	Val	Pro	Gln 20	Asp	Leu	Trp	Ala	Ala 25	Lys	
gcg Ala	tgg Trp	ctg Leu	atc Ile 30	acg Thr	gcc Ala	cgc Arg	agc Ser	ctc Leu 35	tac Tyr	ccg Pro	gca Ala	gac Asp	ttt Phe 40	aac Asn	atc Ile	268
cag Gln	tat Tyr	gag Glu 45	atg Met	tac Tyr	acc Thr	atc Ile	gag Glu 50	cgg Arg	aat Asn	gca Ala	gag Glu	cgg Arg 55	acc Thr	gcc Ala	acc Thr	316
gcc Ala	60 Gly ggg	agg Arg	ctg Leu	ctg Leu	tac Tyr	gac Asp 65	atg Met	ttt Phe	gtg Val	aat Asn	ttc Phe 70	cca Pro	gac Asp	cag Gln	ccg Pro	364
gtg Val 75	gtg Val	tgg Trp	aga Arg	gaa Glu	atc Ile 80	agc Ser	att Ile	att Ile	aca Thr	tca Ser 85	gca Ala	tta Leu	agg Arg	aac Asn	gat Asp 90	412
tca Ser	cag Gln	gac Asp	aaa Lys	caa Gln 95	acc Thr	caa Gln	ttt Phe	tta Leu	aga Arg 100	agt Ser	tta Leu	ttt Phe	gaa Glu	act Thr 105	ctt Leu	460
cct Pro	ggt Gly	cgg Arg	gtc Val 110	cag Gln	tgt Cys	gaa Glu	atg Met	tta Leu 115	cta Leu	aag Lys	gtc Val	acg Thr	gaa Glu 120	caa Gln	tgc Cys	508
ttc Phe	aac Asn	acg Thr 125	tta Leu	gaa Glu	cga Arg	tca Ser	gaa Glu 130	atg Met	ttg Leu	ctt Leu	cta Leu	ctt Leu 135	ttg Leu	agg Arg	cgc Arg	556
ttc Phe	cct Pro 140	Glu	acg Thr	gtg Val	gtg Val	cag Gln 145	cat His	gly ggg	gtt Val	ggc Gly	ctt Leu 150	ggg Gly	gag Glu	gca Ala	cta Leu	604
tta Leu 155	Glu	gct Ala	gaa Glu	act Thr	att Ile 160	Glu	gaa Glu	caa Gln	gaa Glu	tct Ser 165	Pro	gtg Val	aac Asn	tgc Cys	ttt Phe 170	652
aga Arg	aaa Lys	tta Leu	ttt Phe	gtt Val 175	Cys	gat Asp	gtc Val	ctt Leu	cct Pro 180	Leu	ata Ile	att	aac Asn	aac Asn 185	cat His	700
gat Asp	gtt Val	cga Arg	tta Leu 190	Pro	gcc Ala	aat Asn	tta Leu	ttg Leu 195	ı Tyr	aag Lys	tac Tyr	ttg Leu	aac Asn 200	Lys	gca Ala	748
gct Ala	gaa Glu	ttt Phe	туг:	ato : Ile	aat Asr	tat Tyr	gto Val	. Thr	agg Arg	tct Ser	act Thr	caa Glr 215	ı IIE	gaa Glu	aat Asn	796
caç Glr	g cat n His 220	Glr	a ggo n Gly	gco Ala	c caç a Glr	g gat n Asp 225	Thi	tct Ser	gat Asp	tta Lev	a atg 1 Met 230	: Ser	cct Pro	ago Sei	aaa Lys	844
cgt Arg	ago g Sei	c tct	cag Glr	g aag 1 Lys	g tac s Tyı	ata Ile	ata E Ile	a gaa e Glu	a ggg ı Gly	g cto / Lei	g acg ı Thi	g gaa Glu	a aaa 1 Lys	a tca s Sei	tcc Ser	892

235					240				245				250	
												gtt Val 265		940
	_	_	_	_		_	_			-		tat Tyr		988
												aac Asn		1036
												tcc Ser		1084
_	_	_			_		-	_		_		ata Ile		1132
								 _	_		-	ctg Leu 345		1180
												gat Asp		1228
						_						gaa Glu		1276
	_	_		_	_	_						aat Asn		1324
												gtg Val		1372
												tcc Ser 425		1420
												aag Lys		1468
_						_				_	_	atc Ile		1516
												gca Ala		1564

ct Le 47	u GI	g gg n Gl	a tc y Se	c at	t tci e Sei 480	r Glr	g cca	a caq o Gli	g ato n Ilo	c aca e Th: 48!	r Gl	g ca y Gl	g gg n Gl	g ac y Th	c ctg r Leu 490	
ga Gl	g ca u Hi:	t ca s Gl	g ag	g gcg g Ala 499	а ьег	ato 1 Ile	cag Glr	g cto n Lew	g gcg 1 Ala 500	a Thi	g tg Cy	c ca s Hi	c tt s Ph	t gc e Al 50	g cta a Leu 5	1660
999 Gl	g gaq y Gli	g ta ı Ty:	c aga r Arg 510	g Met	g aca Thr	tgt Cys	gaa Glu	a aaa 1 Lys 515	va]	ctt Leu	ga ı Ası	t ttg p Le	g ato u Me 52	t Cy	c tac s Tyr	1708
ato Met	g gta t Val	Let 525	1 Pro	att Ile	caa Gln	gat Asp	gga Gly 530	r Gly	aaa Lys	tco Ser	caç Glı	g gag n Gli 53!	ı Glı	a cco	c tcg Ser	1756
aaa Lys	a gta 5 Val 540	. гу	g cco Pro	aaa Lys	ttt Phe	aga Arg 545	aaa Lys	ggt Gly	tcg Ser	gat Asp	Leu 550	і Гуя	g cto Lev	c ctg 1 Lei	g cct Pro	1804
tgt Cys 555	Thr	ago Ser	aag Lys	gct Ala	atc Ile 560	atg Met	cca Pro	tac Tyr	tgc Cys	ctc Leu 565	cat His	tta Leu	ato Met	j tta : Leu	gcc Ala 570	1852
tgt Cys	ttt Phe	aag Lys	ctt Leu	aga Arg 575	gct Ala	ttc Phe	aca Thr	gac Asp	aac Asn 580	aga Arg	gac Asp	gac Asp	atg Met	gca Ala 585	ttg Leu	1900
Gly aaa	cat His	gtg Val	att Ile 590	gtg Val	ttg Leu	ctt Leu	cag Gln	caa Gln 595	gag Glu	tgg Trp	cca Pro	cgg Arg	ggc Gly 600	Glu	aat Asn	1948
ьеи	PILE	605	гÀг	Ala	Val	Asn	Lys 610	Ile	Cys	Gln	Gln	Gly 615	Asn	Phe	caa Gln	1996
tat Tyr	gag Glu 620	aat Asn	ttt Phe	ttc Phe	aat Asn	tac Tyr 625	gtt Val	aca Thr	aat Asn	att Ile	gat Asp 630	atg Met	ctg Leu	gag Glu	gaa Glu	2044
ttt Phe 635	gcc Ala	tac Tyr	ttg Leu	aga Arg	act Thr 640	cag Gln	gaa Glu	ggt Gly	ggg Gly	aaa Lys 645	att Ile	cat His	ctg Leu	gaa Glu	tta Leu 650	2092
cta Leu	ccc Pro	aat Asn	caa Gln	gga Gly 655	atg Met	ctg Leu	atc Ile	aag Lys	cct Pro 660	tct Ser	agc Ser	cct Pro	ccc Pro	atg Met 665	Gly aaa	2140
шец	ьец	GIN	670	Glu	Phe	tta Leu	Pro	Val 675	Leu	Gln	Pro	Ser	Ile 680	Gln	Thr	2188
gct Ala	voħ	agg Arg 685	cac His	cac His	act of	gta a Val 1	act Thr . 590	cga Arg	ggc Gly	atc Ile	acc Thr	aaa Lys 695	ggc Gly	gtg Val	aag Lys	2236

gag gac ttt cgc ctg gcc atg gag cgc cag gtc tcc cgc tgt gga gag Glu Asp Phe Arg Leu Ala Met Glu Arg Gln Val Ser Arg Cys Gly Glu	2284
700 705 710	
aat ctg atg gtg gtt ctg cac agg ttc tgc att aat gag aag atc ttgAsn Leu Met Val Val Leu His Arg Phe Cys Ile Asn Glu Lys Ile Leu715720	2332
ctc ctt cag act ctg acc tga gt ggagaccttt ccaccagaca cagctcgggc Leu Leu Gln Thr Leu Thr * 735	2385
ctgtgtaatt gtaggagaag acactcagca gtgattgcca tggcacagag ccgtggtcat	2445
tgttgctgtt acaaagaaga aaaccatctg agttctaact ccttggttgc ttaaaagtag	2505
ttcccaagag tctgagaagc tatttctatt tttaagagtc attttttgta atttttgtaa	2565
aacaaaagta ccaatctgtt ttgtaaataa aaatcatcct aaaatttgaa aaaaaaaaaa	2625
<210> 23 <211> 6288 <212> DNA <213> Homo sapiens	
<220> <221> CDS	
<222> (115)(5253)	
<400> 23	60
	60
<400> 23	60 117
<pre><400> 23 aagggagccc cgctcagcgc ggggagcgcc cggcccctc cccgccccat gcgcccgcgg ctctgaagcc tgagcgggc cgggggctgg gcggggccgg ggcccgccgt aggc atg Met</pre>	
<pre></pre>	117
<pre></pre>	117 165
agggagec cgctcagcgc ggggagcgc cggcccctc cccgcccat gcgcccgcgg ctctgaagcc tgagcgggc cgggggctgg gcggggccgg ggcccgccgt aggc atg Met gcg tcc ggg agc cgg tgg cgg ccg aca ccg ccg ccg ctg ctg ttg ctg Ala Ser Gly Ser Arg Trp Arg Pro Thr Pro Pro Pro Leu Leu Leu Leu 5 10 15 ctg ctg ctg gcg ctg gcg gcg cgc gcg gac ggc ctg gag ttc ggc ggc Leu Leu Leu Ala Leu Ala Ala Arg Ala Asp Gly Leu Glu Phe Gly Gly 20 25 30 ggc ccc ggg cag tgg gct cgc tac gcg cgc tgg gcg gcg gcg gcg agc Gly Pro Gly Gln Trp Ala Arg Tyr Ala Arg Trp Ala Gly Ala Ala Ser	117 165 213

ctg gtg gac ggc cgc ctg cgg ctg cgc ttc acg ctt tcg tgc gcc gag 405 Leu Val Asp Gly Arg Leu Arg Leu Arg Phe Thr Leu Ser Cys Ala Glu 90 ceg gee acg etg cag etg gae acg eeg gtg gee gae ege tgg eac 453 Pro Ala Thr Leu Gln Leu Asp Thr Pro Val Ala Asp Asp Arg Trp His 105 atg gtg ctg ctg acc cgc gac gcg cgc cgc acg gcg ctg gcg gtg gac 501 Met Val Leu Leu Thr Arg Asp Ala Arg Arg Thr Ala Leu Ala Val Asp 120 ggc gag gcc cgc gcc gcc gag gtg cgc tcc aag cgg cgc gag atg cag 549 Gly Glu Ala Arg Ala Ala Glu Val Arg Ser Lys Arg Arg Glu Met Gln 135 140 gtg gcc agc gac ctg ttc gtg ggc ggc atc ccg ccc gac gtg cgc ctc 597 Val Ala Ser Asp Leu Phe Val Gly Gly Ile Pro Pro Asp Val Arg Leu 150 tcg gcg ctt acg ctg agc acc gtc aag tac gag ccg ccc ttc cgc ggt 645 Ser Ala Leu Thr Leu Ser Thr Val Lys Tyr Glu Pro Pro Phe Arg Gly 170 ctc ttg gcc aac ctg aag ctg ggc gag cgg ccc ccc gcg ctg ctg ggc 693 Leu Leu Ala Asn Leu Lys Leu Gly Glu Arg Pro Pro Ala Leu Leu Gly age cag gge ctg cgc gcc acc gcc gac ccg ctg tgc gcg ccc gcg 741 Ser Gln Gly Leu Arg Gly Ala Thr Ala Asp Pro Leu Cys Ala Pro Ala 200 cgc aac ccc tgc gcc aac ggc ggc ctc tgc acc gtg ctg gcc ccc ggc 789 Arg Asn Pro Cys Ala Asn Gly Gly Leu Cys Thr Val Leu Ala Pro Gly 210 215 gag gtg ggc tgc gac tgc agc cac acg ggc ttc ggc ggc aag ttc tgc 837 Glu Val Gly Cys Asp Cys Ser His Thr Gly Phe Gly Gly Lys Phe Cys 235 age gaa gag gag cac ccc atg gaa ggt ccg gct cac ctg acg tta aac 885 Ser Glu Glu Glu His Pro Met Glu Gly Pro Ala His Leu Thr Leu Asn 245 age gaa gta ggg tee tta etg tte tee gag ggg gge gee ggg aga gga 933 Ser Glu Val Gly Ser Leu Leu Phe Ser Glu Gly Gly Ala Gly Arg Gly 260 gga gcc ggc gat gtg cac cag cca aca aaa ggc aag gag ttt gtg 981 Gly Ala Gly Asp Val His Gln Pro Thr Lys Gly Lys Glu Glu Phe Val 275 gcg acc ttc aaa ggc aat gag ttc ttc tgc tac gac ctg tca cac aac 1029 Ala Thr Phe Lys Gly Asn Glu Phe Phe Cys Tyr Asp Leu Ser His Asn 290 295

75

80

ccc Pro	atc Ile	cag Gln	agc Ser	agc Ser 310	act Thr	gat Asp	gag Glu	atc Ile	aca Thr 315	ctg Leu	gcc Ala	ttc Phe	cgc Arg	acc Thr 320	ctg Leu	1077
caa Gln	cgc Arg	aac Asn	ggc Gly 325	ctg Leu	atg Met	ctg Leu	cat His	aca Thr 330	ggc Gly	aag Lys	tcg Ser	gcc Ala	gac Asp 335	tac Tyr	gtc Val	1125
aac Asn	ctg Leu	tcc Ser 340	ctc Leu	aag Lys	tct Ser	ggg Gly	gct Ala 345	gtc Val	tgg Trp	ctg Leu	gtc Val	atc Ile 350	aac Asn	cta Leu	ggc Gly	1173
tca Ser	ggt Gly 355	gcc Ala	ttc Phe	gag Glu	gcc Ala	ctt Leu 360	gtg Val	gaa Glu	ccc Pro	gtc Val	aat Asn 365	ggc Gly	aag Lys	ttc Phe	aac Asn	1221
gac Asp 370	aac Asn	gcc Ala	tgg Trp	cac His	gac Asp 375	gtc Val	cgg Arg	gtc Val	acc Thr	cga Arg 380	aac Asn	ctg Leu	cgc Arg	cag Gln	cac His 385	1269
gca Ala	ggg Gly	att Ile	gga Gly	cac His 390	gct Ala	atg Met	gta Val	aac Asn	aaa Lys 395	ctg Leu	cat His	tat Tyr	ctg Leu	gtg Val 400	acc Thr	1317
atc Ile	tcg Ser	gtg Val	gac Asp 405	Gly	atc Ile	ctg Leu	acc Thr	acc Thr 410	aca Thr	ggc Gly	tac Tyr	acg Thr	cag Gln 415	gag Glu	gat Asp	1365
tac Tyr	acc Thr	atg Met 420	ctg Leu	ggc Gly	tct Ser	gat Asp	gac Asp 425	ttc Phe	ttc Phe	tac Tyr	att Ile	999 Gly 430	ggc Gly	agc Ser	ccc Pro	1413
aac Asn	aca Thr 435	gct Ala	gac Asp	ctg Leu	ccg Pro	ggc Gly 440	tcg Ser	ccc Pro	gtc Val	agc Ser	aac Asn 445	aac Asn	ttc Phe	atg Met	ggc Gly	1461
tgc Cys 450	ctc Leu	aag Lys	gac Asp	gtg Val	gtc Val 455	tat Tyr	aag Lys	aac Asn	aat Asn	gac Asp 460	ttc Phe	aaa Lys	ttg Leu	gaa Glu	cta Leu 465	1509
tcc Ser	cgc Arg	ctg Leu	gca Ala	aag Lys 470	gaa Glu	ggg ggg	gac Asp	ccc Pro	aag Lys 475	atg Met	aag Lys	ctg Leu	cag Gln	999 Gly 480	gac Asp	1557
ttg Leu	tca Ser	ttc Phe	cgc Arg 485	tgt Cys	gag Glu	gat Asp	gtg Val	gct Ala 490	gcc Ala	ctg Leu	gac Asp	cct Pro	gtg Val 495	acc Thr	ttt Phe	1605
gag Glu	agt Ser	ccc Pro 500	gag Glu	gcc Ala	ttt Phe	gtg Val	gcg Ala 505	ctg Leu	ccc Pro	cgc Arg	tgg Trp	agc Ser 510	gct Ala	aag Lys	cgc Arg	1653
act Thr	ggc Gly 515	Ser	atc Ile	tcc Ser	cta Leu	gac Asp 520	Phe	cgc Arg	acc Thr	acc Thr	gag Glu 525	Pro	aat Asn	Gly	ctg Leu	1701

			agc Ser													1749
			cag Gln													1797
			ctt Leu 565													1845
			cgc Arg													1893
			Gly 999													1941
_		_	gcc Ala			_	_									1989
_		_	ggc Gly									_	-			2037
			gtg Val 645			_	_			_						2085
			ctc Leu													2133
			cag Gln													2181
			cag Gln													2229
_	_		tgg Trp		_			_	_							2277
			gtc Val 725	_		_		_	_	_						2325
			atg Met													2373
gag	gat	gtg	tcc	ctg	cgt	ttc	atg	tcc	cag	cgg	gcc	tac	gga	ctc	atg	2421

Glu	Asp 755	Val	Ser	Leu	Arg	Phe 760	Met	Ser	Gln	Arg	Ala 765	Tyr	Gly	Leu	Met	
		acc Thr														2469
-		Gly ggg	_	_	_			_			_	_	_	_	_	2517
	_	gca Ala		_				_	_	_					_	2565
		gac Asp 820		_			_				_		_		_	2613
		cag Gln														2661
	-	cat His	_		_							_			_	2709
		cgg Arg														2757
	_	Gly 999						_			_	_	_	_	_	2805
_		gac Asp 900				_				_	_			_	-	2853
		gtg Val														2901
		gcc Ala														2949
	_	acc Thr	_	_						_			_			2997
		gac Asp														3045
		gac Asp														3093

990 985 980 aaa cca gtc aat gac aac cag tgg cac aac gtg gtg gtg tcc agg gac 3141 Lys Pro Val Asn Asp Asn Gln Trp His Asn Val Val Val Ser Arg Asp 1000 995 cca ggc aac gtg cac acg ctc aag att gac tcc cgc act gtc acg cag 3189 Pro Gly Asn Val His Thr Leu Lys Ile Asp Ser Arg Thr Val Thr Gln 1020 1015 1010 cac tcc aat ggc gcc cga aac ctc gat ctc aaa ggg gag ttg tac att 3237 His Ser Asn Gly Ala Arg Asn Leu Asp Leu Lys Gly Glu Leu Tyr Ile 1030 ggc ggt ctg agc aag aat atg ttc agc aac ctg ccc aag ctg gtg gcc 3285 Gly Gly Leu Ser Lys Asn Met Phe Ser Asn Leu Pro Lys Leu Val Ala tee egg gat gge ttt eag gge tge etg gee tea gtg gae ete aac gga 3333 Ser Arg Asp Gly Phe Gln Gly Cys Leu Ala Ser Val Asp Leu Asn Gly cgt ctc cca gac ctc atc gcc gac gcc ctg cac cgc att ggg cag gtg 3381 Arg Leu Pro Asp Leu Ile Ala Asp Ala Leu His Arg Ile Gly Gln Val gag agg ggc tgt gat ggc ccc agc acc acc tgc act gaa gag tcc tgt 3429 Glu Arg Gly Cys Asp Gly Pro Ser Thr Thr Cys Thr Glu Glu Ser Cys 1095 gcc aac cag ggc gtc tgc ttg cag cag tgg gat ggc ttc acc tgc gac 3477 Ala Asn Gln Gly Val Cys Leu Gln Gln Trp Asp Gly Phe Thr Cys Asp 1115 1110 tgc acc atg act tcc tat gga ggc cct gtc tgc aat gat ccc ggg acc 3525 Cys Thr Met Thr Ser Tyr Gly Gly Pro Val Cys Asn Asp Pro Gly Thr 1130 1125 aca tac atc ttt ggg aag ggg gga gcg ctc atc acc tac acg tgg ccc 3573 Thr Tyr Ile Phe Gly Lys Gly Gly Ala Leu Ile Thr Tyr Thr Trp Pro 1150 1145 1140 ccc aat gac agg ccc agc acg agg atg gat cgc ctg gcc gtg ggc ttc 3621 Pro Asn Asp Arg Pro Ser Thr Arg Met Asp Arg Leu Ala Val Gly Phe 1165 1160 1155 age ace cae cag egg age get gtg etg gtg egg gtg gae age gee tee 3669 Ser Thr His Gln Arg Ser Ala Val Leu Val Arg Val Asp Ser Ala Ser 1185 1175 1170 ggc ctt gga gac tac ctg cag ctg cac atc gac cag ggc acc gtg ggg 3717

3765

Gly Leu Gly Asp Tyr Leu Gln Leu His Ile Asp Gln Gly Thr Val Gly

gtg atc ttt aac gtg ggc acg gac gac att acc atc gac gag ccc aac

Val Ile Phe Asn Val Gly Thr Asp Asp Ile Thr Ile Asp Glu Pro Asn 1210

1190

	ata Ile					Lys					Arg					3813
ggc Gly	ggc Gly 1235	aac Asn	gcc Ala	acc Thr	Leu	cag Gln 1240	gtg Val	gac Asp	agc Ser	Trp	ccg Pro 1245	gtc Val	aac Asn	gag Glu	cgg Arg	3861
	ccg Pro			Asn					Arg					Arg		3909
	atc Ile		Tyr					Val					Leu			3957
	ggc Gly	Arg					Phe					Ala				4005
	ggc Gly					Arg					Gln					4053
Tyr	tac Tyr 1315				Lys					Ala						4101
aat Asn 1330	gtg Val	cgg Arg	act Thr	Glu	ggt Gly 1335	cac His	ctg Leu	cgc Arg	Leu	gtg Val L340	ggg Gly	gag Glu	gly ggg	Pro	tcc Ser 1345	4149
	ctg Leu		Ser					Ala					Āla			4197
	acc Thr	Thr					Thr					Thr				4245
cgc Arg	cgg Arg	ggc Gly 1380	cgc Arg	tcc Ser	ccc Pro	Thr	ctg Leu .385	agg Arg	gac Asp	agc Ser	Thr	acc Thr 390	cag Gln	aac Asn	aca Thr	4293
Asp	gac Asp 1395				Ala					Pro						4341
	gag Glu			Glu					Gly					Pro		4389
	acg Thr		Asp					Pro					Arg			4437

	gtg Val	Pro					Phe					Thr				4485
	acc Thr					Pro					Arg					4533
Gly	ggc Gly 1475				Ala					Ser						4581
	gag Glu			Gly					Glu					Ser		4629
	ccc Pro		Asp					Tyr					Leu			4677
	cgc Arg	Thr					Pro				_	Pro				4725
	agg Arg					Thr					Val					4773
Ser	gcc Ala 1555				Asn					Lys						4821
	ctt Leu			Leu					Pro					Ala		4869
	tcc Ser		Glu					Pro					Gly			4917
	gcc Ala	Pro					Leu					Pro				4965
	gag Glu					Gly					Ile					5013
Ser	acc Thr 1635				Val					Ala						5061
	ctc Leu			Leu					Lys					Asp		5109
ggc	tcc	tac	cag	gtg	gac	cag	agc	cga	aac	tac	atc	agt	aac	tcg	gcc	5157

Gly Ser Tyr Gln Val Asp Gln Ser Arg Asn Tyr Ile Ser Asn Ser Ala 1670 1675 1680	
cag agc aat ggg gcg gtg gtg aaa gag aag gcc ccg gct gcc ccc aag Gln Ser Asn Gly Ala Val Val Lys Glu Lys Ala Pro Ala Ala Pro Lys 1685 1690 1695	5205
acg ccc agc aag gcc aag aag aac aaa gac aag gag tat tat gtc tga Thr Pro Ser Lys Ala Lys Lys Asn Lys Asp Lys Glu Tyr Tyr Val * 1700 1705 1710	5253
gccccggca ctgcgcccca ctgccagctg cccctcctgg gagggcccgg gaggagggtg	5313
ccaccetete cetgecaggg geetggggae cetetecetg getgeeteag gettetetta	5373
cgaagaggaa acgcaaaaaa agaaaaggaa aaaccccgtg ctcgccccct tcctcctgcc	5433
gtccactgcg cggcctcgtc agtcccgggg ctgactgtcc ctctcagctc tgcgcctgcc	5493
aggcagggca cgtgctcaca gccctgggtt gatttatttt tttaaggggg tagtttatt	5553
ttggtggggt tgggtgggaa ggaaggctgg gggttttgta aagtgtccac tgctcgtcct	5613
gttaattttc ctcaattttt cttcttcttc cttctgtccc tcctgccttc cttctcttcc	5673
caagecetee aateceeate ecaggettge tgtgteteae tgteeecace etectteeet	5733
acttettttt ttgtgtgtet ggtttetece tteettteet	5793
ggtgggagaa gggcgggagg gtgggcccga gtggcccagt gggtgggtgg ggtggggtgg	5853
ggcaagtgcc ccaactcccc tcaccaggag aggcacctgc ttggtgccgc ccagggaagg	5913
ggctcaggcc tgacggaagg cctgttctgt gtgtgccgcc gggcgacgtg cattgatggg	5973
gaagctgctg gaggagcagg ggtgggggt gggagggagg ggaaaggcaa atgcagatat	6033
atattacaga caaatactct agattccacg agcagcagcc tgtggcaccc gctgggcgcg	6093
ggcagcaggg aagagggagc aaggcattgt ccacagactg ctggggtcac ttctttgccc	6153
acgggctccc tgctccccca gttttttttc tctctttgtt aacaaatgtg tctgagtctt	6213
ggaaaacacc ccaaccccgg aaatgtgtgg gaaaaagaaa acaaaaactt tccaaattcc	6273
aaaaaaaaaa aaaaa	6288

<210> 24

<211> 1016 <212> DNA

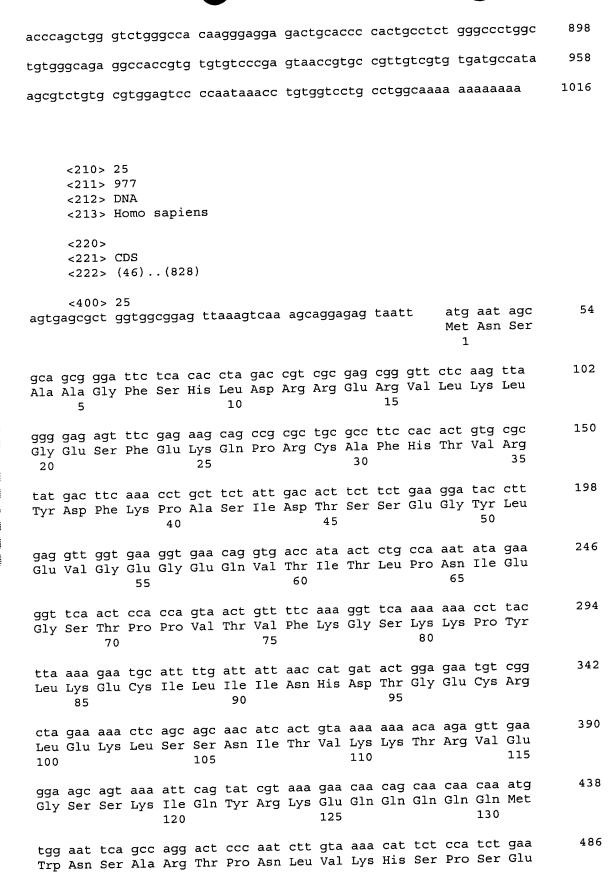
<213> Homo sapiens

<220>

<221> CDS

<222> (254)..(724)

<400> 24 ttaatacgat gactataggg aatttggccc tcgagcaaga attcggcacg aggctggctc	60
tgactettea eccatettea eccaggetgg eccetttggt gaaactacaa etceeagggg	120
tetgtgegeg agaaggeagg egggttttte taceggaagt eegetetage tetgggeeet	180
acaactgcac cctgagccgg agctgcccag tcgccgcggg accggggccg ctggggtctg	240
gacgggggtc gcc atg atc cgc ttt atc ctc atc cag aac cgg gca ggc Met Ile Arg Phe Ile Leu Ile Gln Asn Arg Ala Gly 1 5 10	289
aag acg cgc ctg gcc aag tgg tac atg cag ttt gat gat gat gag aaa Lys Thr Arg Leu Ala Lys Trp Tyr Met Gln Phe Asp Asp Asp Glu Lys 15 20 25	337
cag aag ctg atc gag gag gtg cat gcc gtg gtc acc gtc cga gac gcc Gln Lys Leu Ile Glu Glu Val His Ala Val Val Thr Val Arg Asp Ala 30 35 40	385
aaa cac acc aac ttt gtg gag gtc ctg gca agc tcc gtt gct gac agc Lys His Thr Asn Phe Val Glu Val Leu Ala Ser Ser Val Ala Asp Ser 45 50 55 60	433
ctc tct gtt ctg cag ttc cgg aac ttt aag atc att tac cgc cgc tat Leu Ser Val Leu Gln Phe Arg Asn Phe Lys Ile Ile Tyr Arg Arg Tyr 65 70 75	481
gct ggc ctc tac ttc tgc atc tgt gtg gat gtc aat gac aac aac ctg Ala Gly Leu Tyr Phe Cys Ile Cys Val Asp Val Asn Asp Asn Asn Leu 80 85 90	529
gct tac ctg gag gcc att cac aac ttc gtg gag gtc tta aac gaa tat Ala Tyr Leu Glu Ala Ile His Asn Phe Val Glu Val Leu Asn Glu Tyr 95 100 105	577
ttc cac aat gtc tgt gaa ctg gac ctg gtg ttc aac ttc tac aag gtt Phe His Asn Val Cys Glu Leu Asp Leu Val Phe Asn Phe Tyr Lys Val 110 115 120	625
tac acg gtc gtg gac gag atg ttc ctg gct ggc gaa atc cga gag acc Tyr Thr Val Val Asp Glu Met Phe Leu Ala Gly Glu Ile Arg Glu Thr 125 130 135 140	673
agc cag acg aag gtg ctg aaa cag ctg ctg atg cta cag tcc ctg gag Ser Gln Thr Lys Val Leu Lys Gln Leu Leu Met Leu Gln Ser Leu Glu 145 150 155	721
tga gggc aggcgagccc caccccggcc ccggcccctc ctggactcgc ctgctcgctt *	778
ccccttccca ggcccgtggc caacccagca gtccttccct cagctgccta ggaggaaggg	838



145 140 135 gat aag atg tcc cca gca tct cca ata gat gat atc gaa aga gaa ctg 534 Asp Lys Met Ser Pro Ala Ser Pro Ile Asp Asp Ile Glu Arg Glu Leu 155 150 582 aag gca gaa gct agt cta atg gac cag atg agt agt tgt gat agt tca Lys Ala Glu Ala Ser Leu Met Asp Gln Met Ser Ser Cys Asp Ser Ser 175 170 165 tca gat tcc aaa agt tca tca tct tca agt agt gag gat agt tct agt 630 Ser Asp Ser Lys Ser Ser Ser Ser Ser Ser Ser Glu Asp Ser Ser Ser 185 180 gac tca gaa gat gaa gat tgc aaa tcc tct act tct gat aca ggg aat 678 Asp Ser Glu Asp Glu Asp Cys Lys Ser Ser Thr Ser Asp Thr Gly Asn 210 200 tgt gtc tca gga cat cct acc atg aca cag tac agg att cct gat ata 726 Cys Val Ser Gly His Pro Thr Met Thr Gln Tyr Arg Ile Pro Asp Ile 215 774 gat gcc agt cat aat aga ttt cga gac aac agt ggc ctt ctg atg aat Asp Ala Ser His Asn Arg Phe Arg Asp Asn Ser Gly Leu Leu Met Asn 235 act tta aga aat gat ttg cag ctg agt gaa tca gga agt gac agt gat 822 Thr Leu Arg Asn Asp Leu Gln Leu Ser Glu Ser Gly Ser Asp Ser Asp 255 250 245 gac tga agaaatattt agctataaat aaaaatttat acagcatgta taatttattt 878 260 tgtattaaca ataaaaattc ctaagactga gggaaatatg tcttaacttt tgatgataaa 938 977 agaaattaaa tttgattcag aaatttcaaa aaaaaaaaa <210> 26 <211> 3185 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (37)..(1257) <400> 26 accoeggegg coetgecege coetecetee ageate atg gee age eea aga acc 54 Met Ala Ser Pro Arg Thr 1

agg aag gtt ctt aaa gaa gtc agg gtg cag gat gag aac aac gtt tgt Arg Lys Val Leu Lys Glu Val Arg Val Gln Asp Glu Asn Asn Val Cys

20 15 10 150 ttt gag tgt ggc gcg ttc aat cct cag tgg gtc agt gtg acc tac ggc Phe Glu Cys Gly Ala Phe Asn Pro Gln Trp Val Ser Val Thr Tyr Gly 30 198 atc tgg atc tgc ctg gag tgc tcg ggg aga cac cgc ggg ctt ggg gtt Ile Trp Ile Cys Leu Glu Cys Ser Gly Arg His Arg Gly Leu Gly Val 246 cac ctc agc ttt gtg cgc tct gtt act atg gac aag tgg aag gac att His Leu Ser Phe Val Arg Ser Val Thr Met Asp Lys Trp Lys Asp Ile 60 55 294 gag ctt gag aag atg aaa gct ggt ggg aat gct aag ttc cga gag ttc Glu Leu Glu Lys Met Lys Ala Gly Gly Asn Ala Lys Phe Arg Glu Phe ctg gag tct cag gag gat tac gat cct tgc tgg tcc ttg cag gag aag 342 Leu Glu Ser Gln Glu Asp Tyr Asp Pro Cys Trp Ser Leu Gln Glu Lys 390 tac aac agc aga gcc gcg gcc ctc ttt agg gat aag gtg gtc gct ctg Tyr Asn Ser Arg Ala Ala Ala Leu Phe Arg Asp Lys Val Val Ala Leu 438 gcc gaa ggc aga gag tgg tct ctg gag tca tca cct gcc cag aac tgg Ala Glu Gly Arg Glu Trp Ser Leu Glu Ser Ser Pro Ala Gln Asn Trp 125 120 486 acc cca cct cag ccc agg acg ctg ccg tcc atg gtg cac cga gtc tct Thr Pro Pro Gln Pro Arg Thr Leu Pro Ser Met Val His Arg Val Ser 140 145 135 ggc cag ccg cag agt gtg acc gcc tcc tcg gac aag gct ttt gaa gac 534 Gly Gln Pro Gln Ser Val Thr Ala Ser Ser Asp Lys Ala Phe Glu Asp 155 160 tgg ctg aat gat gac ctc ggc tcc tat caa ggg gcc cag ggg aat cgc 582 Trp Leu Asn Asp Asp Leu Gly Ser Tyr Gln Gly Ala Gln Gly Asn Arg 175 170 630 tac gtg ggg ttt ggg aac acg cca ccg cct cag aag aaa gaa gat gac Tyr Val Gly Phe Gly Asn Thr Pro Pro Pro Gln Lys Lys Glu Asp Asp 190 185 ttc ctc aac aac gcc atg tcc tcc ctg tac tcg ggc tgg agc agc ttc 678 Phe Leu Asn Asn Ala Met Ser Ser Leu Tyr Ser Gly Trp Ser Ser Phe 205 200 acc act gga gcc agc cgg ttt gcc tcg gca gcc aag gag ggc gct aca 726 Thr Thr Gly Ala Ser Arg Phe Ala Ser Ala Ala Lys Glu Gly Ala Thr 230 220 774 aag ttt gga tcc caa gcg agt cag aag gcg tcc gag ctg ggc cac agc Lys Phe Gly Ser Gln Ala Ser Gln Lys Ala Ser Glu Leu Gly His Ser

240

ctg aac gag aac gtc ctc aag cct gcg Leu Asn Glu Asn Val Leu Lys Pro Ala 250 255	
aag att ttt gat gat gtc tcc agt ggg Lys Ile Phe Asp Asp Val Ser Ser Gly 265 270	
gtc cag gga gtc ggt agt aag gga tgg Val Gln Gly Val Gly Ser Lys Gly Trp 280 285	
tcg ggg aaa gca gag ggc ccc ttg gac Ser Gly Lys Ala Glu Gly Pro Leu Asp 295 300	
tat cag aac agc ggt ctg gac cac ttc Tyr Gln Asn Ser Gly Leu Asp His Phe 315	
agc ttc tgg gag acc ttt gga agt gct Ser Phe Trp Glu Thr Phe Gly Ser Ala 330 335	
tcc ccg agc agc gac agc tgg acg tgc Ser Pro Ser Ser Asp Ser Trp Thr Cys 345 350	
agg agc tcg gac agc tgg gag gtg tgg Arg Ser Ser Asp Ser Trp Glu Val Trp 360 365	
aac agc aac agc gac ggc ggg gag ggc Asn Ser Asn Ser Asp Gly Gly Glu Gly 375 380	
gtg ccg ccg gcc gtg ccc act gat gat Val Pro Pro Ala Val Pro Thr Asp Asp 395	
tag ggcc cactgcgccc ccgtccccag cgcc *	cccggg cgacttcgtg tttgcactct 1311
gccctcgtcg ttcctcctcc ttccatttga cc	caagaatc agcaactgca gtgtgaggac 1371
agegtetegg gaggeaggae eetagggaga ee	cgggtgtg cgccgcctgc gcgtggggag 1431
tetteggtge gtgggggegg ettgetgtee ag	cctgtgtg ggggccgtcc cgtcccacac 1491
teceetgge attettggae teaaggeegg gg	ctctgcgt ggcttgctgg gaggtgggct 1551
gcagcacaga ggcctgtgac tgcgttccag cg	gccagttc actacgcagt atctctgggg 1611
cetgggacca gccacgtgcc gagetgtcag eg	acgtgagg tgtcccttct cgttgagata 1671



<210> 27 <211> 1850

<212> DNA <213> Homo sapiens <220> <221> CDS <222> (596)..(1840) <400> 27 60 tttttgtatg caccacgggc ggcggtggtc ggtgcgggag gagggagggg agcttgcggg cccgagaggg ggcgacggcg gcggcggtgg cctgaggagg cccgagcggc ggcggtggcg 120 gcgaaggccg aggcgtctag gtgtttttgg aagagctgca gccctcttct cacagatgag 180 ctacgaggag atgatgacac tgactgagca gcacctggag tctcagaacg tcaccaaagg 240 tgcccgccac aagatagccc tgagcatcca gaagctgcgt gagagacaga gcgtcctcaa 300 gtccctagag aaggatgtgc tggaaggcgg gaacctacga aacgctctgc aggagctgca 360 gcagatcatc atcactccca tcaaggccta cagtgtcctc caggccaccg tggctgccgc 420 caccaccacc cctactgcca aggatggggc cccgggggaa ccaccgctgc caggtgctga 480 540 gectececta geceaeceeg geacagaeaa aggeaecgag gecagageeg ggaecatgtg acggcgctgg ccctcgccac cgccgtcccc cgaccctggc cccaggcccg gcacc atg 598 Met atg ttc cga gac cag gtg ggc atc ctc gct ggc tgg ttc aaa ggc tgg 646 Met Phe Arg Asp Gln Val Gly Ile Leu Ala Gly Trp Phe Lys Gly Trp aat gag tgt gag cag aca gtg gcc ctc ctg tca ctt ccg aaa cgg gtc 694 Asn Glu Cys Glu Gln Thr Val Ala Leu Leu Ser Leu Pro Lys Arg Val 25 20 acc cgt acc cag gcc cgc ttc ctg cag ctc tgc ctg gag cac tca ctg 742 Thr Arg Thr Gln Ala Arg Phe Leu Gln Leu Cys Leu Glu His Ser Leu 40 35 gcg gac tgc aat gac atc cac ctg ctg gag tcg gag gcc aac agt gct 790 Ala Asp Cys Asn Asp Ile His Leu Leu Glu Ser Glu Ala Asn Ser Ala 60 55 50 gcc atc gtc agc cag tgg cag cag gag tcc aaa gag aag gtg gtg tcc 838 Ala Ile Val Ser Gln Trp Gln Gln Glu Ser Lys Glu Lys Val Val Ser 70 886 ctc ctg ctg tcc cac ctt ccc ctg ctt cag cca ggc aac aca gag gcc Leu Leu Ser His Leu Pro Leu Leu Gln Pro Gly Asn Thr Glu Ala 85

934

aag tcg gag tac atg agg cta ctg cag aaa gtg ctg gcc tac tca atc

Lys Ser Glu Tyr Met Arg Leu Leu Gln Lys Val Leu Ala Tyr Ser Ile

105

gag Glu	agc Ser 115	aat Asn	gct Ala	ttc Phe	atc Ile	gag Glu 120	gag Glu	agt Ser	cgc Arg	cag Gln	ctg Leu 125	ctt Leu	tcc Ser	tat Tyr	gcc Ala	982
ctc Leu 130	atc Ile	cac His	cca Pro	gcc Ala	acc Thr 135	aca Thr	ctg Leu	gag Glu	gac Asp	cgc Arg 140	aac Asn	gca Ala	ctg Leu	gcc Ala	ctc Leu 145	1030
tgg Trp	ctg Leu	agc Ser	cac His	ctg Leu 150	gaa Glu	gag Glu	cgg Arg	ttg Leu	gct Ala 155	agt Ser	ggc Gly	ttc Phe	cgc Arg	tcc Ser 160	cgg Arg	1078
cca Pro	gag Glu	ccc Pro	tcc Ser 165	tac Tyr	cat His	tca Ser	cgt Arg	caa Gln 170	ggc Gly	tca Ser	gat Asp	gag Glu	tgg Trp 175	gly ggg	ggc Gly	1126
cct Pro	gca Ala	gag Glu 180	cta Leu	ggc Gly	cct Pro	gly ggg	gag Glu 185	gca Ala	ggg Gly	cca Pro	ggc Gly	tgg Trp 190	cag Gln	gac Asp	aag Lys	1174
cca Pro	ccc Pro 195	cgg Arg	gaa Glu	aat Asn	gga Gly	cac His 200	gtg Val	ccc Pro	ttc Phe	cac His	cca Pro 205	tcc Ser	agc Ser	tca Ser	gtg Val	1222
ccg Pro 210	cca Pro	gcc Ala	atc Ile	aac Asn	agt Ser 215	att Ile	ggg Gly	agc Ser	aat Asn	gca Ala 220	aac Asn	aca Thr	ggt Gly	ctc Leu	ccc Pro 225	1270
tgc Cys	caa Gln	atc Ile	cac His	cct Pro 230	agc Ser	cca Pro	ctg Leu	aag Lys	cgc Arg 235	tcc Ser	atg Met	tca Ser	ctc Leu	atc Ile 240	cct Pro	1318
aca Thr	agc Ser	ccc Pro	cag Gln 245	gtc Val	cct Pro	ggt Gly	gag Glu	tgg Trp 250	Pro	agt Ser	cca Pro	gag Glu	gag Glu 255	ctt Leu	gly aaa	1366
gcc Ala	cgg Arg	gct Ala 260	gct Ala	ttt Phe	acc Thr	acg Thr	ccc Pro 265	Asp	cac His	gca Ala	cct Pro	ctc Leu 270	tcg Ser	ccc Pro	cag Gln	1414
agc Ser	agc Ser 275	Val	gcc Ala	tcc Ser	tct Ser	ggc Gly 280	Ser	gag Glu	cag Gln	aca Thr	gag Glu 285	Glu	cag Gln	ggc Gly	tcc Ser	1462
ago Ser 290	Arg	aac Asn	acc Thr	ttc Phe	cag Gln 295	Glu	gat Asp	ggc Gly	agt Ser	ggc Gly 300	Met	aaa Lys	gat Asp	gtg Val	ccc Pro 305	1510
tca Ser	tgg Trp	cto Leu	aag Lys	ago Ser 310	Leu	cgt Arg	ttg Leu	cac His	aag Lys 315	Tyr	gca Ala	gcc Ala	ctc Leu	tto Phe 320	tca Ser	1558
caç Glr	g atg n Met	ago Ser	tac Tyr 325	Glu	gag Glu	atg Met	atg Met	aca Thr	Leu	act Thr	gag Glu	g cag i Glr	cac His	Leu	gag Glu	1606

tct cag aac gtc acc aaa ggt gcc cgc cac aag ata gcc ctg agc atc Ser Gln Asn Val Thr Lys Gly Ala Arg His Lys Ile Ala Leu Ser Ile 340 345 350	1654
cag aag ctg cgt gag aga cag agc gtc ctc aag tcc cta gag aag gat Gln Lys Leu Arg Glu Arg Gln Ser Val Leu Lys Ser Leu Glu Lys Asp 355 360 365	1702
gtg ctg gaa ggc ggg aac cta cga aac gct ctg cag gag ctg cag cag Val Leu Glu Gly Gly Asn Leu Arg Asn Ala Leu Gln Glu Leu Gln Gln 370 375 380 385	1750
atc atc atc ccc atc aag gcc tac agt gtc ctc cag gcc acc gtg Ile Ile Ile Thr Pro Ile Lys Ala Tyr Ser Val Leu Gln Ala Thr Val 390 395 400	1798
gct gcc gcc acc ctg tat tgc ggc cgc tct aga gga tcc tag cttcgta Ala Ala Ala Thr Leu Tyr Cys Gly Arg Ser Arg Gly Ser * 405 410 415	1847
cgc	1850
<210> 28	
<211> 4781 <212> DNA <213> Homo sapiens	
(213) 1101110 2421-1-1	
<220>	
<220> <221> CDS <222> (813)(3635)	
<221> CDS <222> (813)(3635)	60
<221> CDS <222> (813)(3635) <400> 28 tttgctgtgg aattcccggg tcgacccacg cgtccgcgga cgcgtgggtt tagaggtaag	60 120
<pre><221> CDS <222> (813)(3635) <400> 28 tttgctgtgg aattcccggg tcgacccacg cgtccgcgga cgcgtgggtt tagaggtaag tttgcctact ttgtcgtcta gtgggtaaaa ttttgcggag agcgttggat ctgggaagcg</pre>	
<pre><221> CDS <222> (813)(3635) <400> 28 tttgctgtgg aattcccggg tcgacccacg cgtccgcgga cgcgtgggtt tagaggtaag tttgcctact ttgtcgtcta gtgggtaaaa ttttgcggag agcgttggat ctgggaagcg ggatagggat ggatgggttc atttgagagc cacggcttaa agcggttgcg atcaggatgg</pre>	120
<pre><221> CDS <222> (813)(3635) <400> 28 tttgctgtgg aattcccggg tcgacccacg cgtccgcgga cgcgtgggtt tagaggtaag tttgcctact ttgtcgtcta gtgggtaaaa ttttgcggag agcgttggat ctgggaagcg ggatagggat ggatgggttc atttgagagc cacggcttaa agcggttgcg atcaggatgg gacacaggtt tgtttgggga caacaaagat ggcatttgtg agtgttttga agcaacccgt</pre>	120 180
<pre> <221> CDS <222> (813)(3635) <400> 28 tttgctgtgg aattcccggg tcgacccacg cgtccgcgga cgcgtgggtt tagaggtaag tttgcctact ttgtcgtcta gtgggtaaaa ttttgcggag agcgttggat ctgggaagcg ggatagggat ggatgggttc atttgagagc cacggcttaa agcggttgcg atcaggatgg gacacaggtt tgtttgggga caacaaagat ggcatttgtg agtgttttga agcaacccgt actgattaca tctttctccc ttgtgttcct tttatcccag gtttgaattt tctcggagaa </pre>	120 180 240
<pre> <221> CDS <222> (813)(3635) </pre> <pre> <400> 28 tttgctgtgg aattcccggg tcgacccacg cgtccgcgga cgcgtgggtt tagaggtaag tttgcctact ttgtcgtcta gtgggtaaaa ttttgcggag agcgttggat ctgggaagcg ggatagggat ggatgggttc atttgagagc cacggcttaa agcggttgcg atcaggatgg gacacaggtt tgtttgggga caacaaagat ggcatttgtg agtgtttga agcaacccgt actgattaca tctttctccc ttgtgttcct tttatcccag gtttgaattt tctcggagaa agacaggccg gccacgagga aaacagaaac aagccgcagc aacatctaag cccttgaaag</pre>	120 180 240 300
<pre> <221> CDS <222> (813)(3635) <400> 28 tttgctgtgg aattcccggg tcgacccacg cgtccgcgga cgcgtgggtt tagaggtaag tttgcctact ttgtcgtcta gtgggtaaaa ttttgcggag agcgttggat ctgggaagcg ggatagggat ggatgggttc atttgagagc cacggcttaa agcggttgcg atcaggatgg gacacaggtt tgtttgggga caacaaagat ggcatttgtg agtgttttga agcaacccgt actgattaca tctttctccc ttgtgttcct tttatcccag gtttgaattt tctcggagaa </pre>	120 180 240 300 360
<pre><221> CDS <222> (813)(3635) </pre> <pre> <400> 28 tttgctgtgg aattcccggg tcgacccacg cgtccgcgga cgcgtgggtt tagaggtaag tttgcctact ttgtcgtcta gtgggtaaaa ttttgcggag agcgttggat ctgggaagcg ggatagggat ggatgggttc atttgagagc cacggcttaa agcggttgcg atcaggatgg gacacaggtt tgtttgggga caacaaagat ggcatttgtg agtgtttga agcaacccgt actgattaca tctttctccc ttgtgttcct tttatcccag gtttgaattt tctcggagaa agacaggccg gccacgagga aaacagaaac aagccgcagc aacatctaag cccttgaaag gatcctgaga gagggggaa agggaaaaca gcagccacca gcccaaccac ttgtgtcttc</pre>	120 180 240 300 360 420
<pre> <221> CDS <222> (813)(3635) <400> 28 tttgctgtgg aattcccggg tcgacccacg cgtccgcgga cgcgtgggtt tagaggtaag tttgcctact ttgtcgtcta gtgggtaaaa ttttgcggag agcgttggat ctgggaagcg ggatagggat ggatgggttc atttgagagc cacggcttaa agcggttgcg atcaggatgg gacacaggtt tgtttgggga caacaaagat ggcatttgtg agtgtttga agcaacccgt actgattaca tctttctccc ttgtgttcct tttatcccag gtttgaattt tctcggagaa agacaggccg gccacgagga aaacagaaac aagccgcagc aacatctaag cccttgaaag gatcctgaga gagggggaa agggaaaaca gcagccacca gcccaaccac ttgtgtcttc tgccccttcc cacctatctt gcccaccca ccagcccacg ctgcttgga cttgaaatct </pre>	120 180 240 300 360 420 480

ttttaaaaga ctttttgatc caatgaggcc ccctaaataa ttgagttttg ggtcctggtt	720
ggttgtttta tttttttcc tccaaaattt taccccctcc cccctgagcc cgaggtgctg	780
acgtcgcaaa aaaattggat aaaaccacca to atg ggt tcg ggt ccc ata gac Met Gly Ser Gly Pro Ile Asp 1 5	833
ccc aaa gaa ctt ctc aag ggc ctg gac agc ttc ctt aac cga gat ggg Pro Lys Glu Leu Lys Gly Leu Asp Ser Phe Leu Asn Arg Asp Gly 10 15 20	881
gaa gtc aaa agt gtg gat ggg att tcc aag atc ttc agt ttg atg aag Glu Val Lys Ser Val Asp Gly Ile Ser Lys Ile Phe Ser Leu Met Lys 25 30 35	929
gaa gca cga aag atg gtg agt cga tgc act tac ttg aac att ctc ctg Glu Ala Arg Lys Met Val Ser Arg Cys Thr Tyr Leu Asn Ile Leu Leu 40 45 50 55	977
cag acc cgt tca cca gaa ata ttg gtc aaa ttt att gac gtt ggc ggc Gln Thr Arg Ser Pro Glu Ile Leu Val Lys Phe Ile Asp Val Gly 60 65 70	1025
tac aaa ctt ctt aac aat tgg ctg acg tat tca aag aca acc aac aac Tyr Lys Leu Leu Asn Asn Trp Leu Thr Tyr Ser Lys Thr Thr Asn Asn 75 80 85	1073
att ccc ctc ctc cag caa att cta ctg acc ctg cag cat cta ccg ctc Ile Pro Leu Leu Gln Gln Ile Leu Leu Thr Leu Gln His Leu Pro Leu 90 95 100	1121
act gta gac cat ctc aag cag aac aac aca gct aaa ctg gtg aag cag Thr Val Asp His Leu Lys Gln Asn Asn Thr Ala Lys Leu Val Lys Gln 105	1169
ctg agc aag tca agt gag gat gaa gag ctc cgg aaa ttg gcc tca gtc Leu Ser Lys Ser Ser Glu Asp Glu Glu Leu Arg Lys Leu Ala Ser Val 120 125 130 135	1217
ctt gtc agc gac tgg atg gct gtc atc cgc tct cag agc agt acc cag Leu Val Ser Asp Trp Met Ala Val Ile Arg Ser Gln Ser Ser Thr Gln 140 145 150	1265
cct gct gag aaa gat aag aag aaa cgt aaa gat gaa gga aaa agt cga Pro Ala Glu Lys Asp Lys Lys Lys Arg Lys Asp Glu Gly Lys Ser Arg 155 160 165	1313
acc acc ctt cct gag cga cct ttg aca gag gtg aag gct gag acc cgg Thr Thr Leu Pro Glu Arg Pro Leu Thr Glu Val Lys Ala Glu Thr Arg 170 175 180	1361
gct gag gag gcc cca gag aag aag agg gag aag ccc aag tct ctt cgc Ala Glu Glu Ala Pro Glu Lys Lys Arg Glu Lys Pro Lys Ser Leu Arg 185	1409
acc aca gca ccc agt cat gcc aag ttc cgt tcc act gga cta gag ctg	1457

Thr 200	Thr	Ala	Pro	Ser	His 205	Ala	Lys	Phe	Arg	Ser 210	Thr	Gly	Leu	Glu	Leu 215	
gag Glu	aca Thr	cca Pro	tcc Ser	ttg Leu 220	gtg Val	cct Pro	gtg Val	aag Lys	aag Lys 225	aat Asn	gcc Ala	agc Ser	aca Thr	gtg Val 230	gtg Val	1505
gtt Val	tct Ser	gac Asp	aag Lys 235	tac Tyr	aac Asn	ctt Leu	aaa Lys	ccc Pro 240	atc Ile	ccc Pro	ctc Leu	aaa Lys	cgt Arg 245	cag Gln	agc Ser	1553
aac Asn	gta Val	gct Ala 250	gct Ala	cca Pro	gga Gly	gat Asp	gcc Ala 255	act Thr	ccc Pro	cct Pro	gca Ala	gag Glu 260	aag Lys	aaa Lys	tac Tyr	1601
aag Lys	cca Pro 265	ctc Leu	aac Asn	aca Thr	aca Thr	cct Pro 270	aat Asn	gcc Ala	acc Thr	aaa Lys	gag Glu 275	atc Ile	aaa Lys	gtg Val	aag Lys	1649
atc Ile 280	atc Ile	ccg Pro	cca Pro	cag Gln	cct Pro 285	atg Met	gag Glu	ggc Gly	ctg Leu	ggc Gly 290	ttt Phe	ctg Leu	gat Asp	gct Ala	ctt Leu 295	1697
aat Asn	tca Ser	gcc Ala	cct Pro	gtt Val 300	cca Pro	ggc Gly	atc Ile	aaa Lys	att Ile 305	aag Lys	aag Lys	aaa Lys	aaa Lys	aaa Lys 310	gta Val	1745
ctg Leu	tca Ser	cct Pro	acg Thr 315	gct Ala	gcc Ala	aag Lys	cca Pro	agc Ser 320	ccc Pro	ttt Phe	gaa Glu	Gly aaa	aaa Lys 325	acg Thr	agc Ser	1793
aca Thr	gaa Glu	cca Pro 330	agc Ser	aca Thr	gcc Ala	aaa Lys	cct Pro 335	tct Ser	tcc Ser	cca Pro	gaa Glu	cca Pro 340	gca Ala	cca Pro	cct Pro	1841
tct Ser	gag Glu 345	gca Ala	atg Met	gac Asp	gca Ala	gac Asp 350	cgt Arg	cca Pro	ggc Gly	acc Thr	ccg Pro 355	gtt Val	ccc Pro	cct Pro	gtt Val	1889
gaa Glu 360	gtc Val	ccg Pro	gag Glu	ctc Leu	atg Met 365	gat Asp	aca Thr	gcc Ala	tct Ser	ttg Leu 370	gag Glu	cca Pro	gga Gly	gct Ala	ctg Leu 375	1937
gat Asp	gcc Ala	aag Lys	cca Pro	gtg Val 380	Glu	agt Ser	cct Pro	gga Gly	gat Asp 385	cct Pro	aac Asn	caa Gln	ctg Leu	acc Thr 390	Arg	1985
aaa Lys	ggc Gly	agg Arg	aag Lys 395	Arg	aaa Lys	agt Ser	gtg Val	aca Thr 400	Trp	cct Pro	gag Glu	gaa Glu	ggc Gly 405	Lys	ctg Leu	2033
aga Arg	gaa Glu	tat Tyr 410	ttc Phe	tat Tyr	ttt Phe	gaa Glu	ttg Leu 415	Asp	gaa Glu	act Thr	gaa Glu	cga Arg 420	Val	aat Asn	gtg Val	2081
aat Asn	aag Lys	atc Ile	aag Lys	gac	ttt Phe	ggt Gly	gag Glu	gcg Ala	gct Ala	aag Lys	cga Arg	gag Glu	ata Ile	ctg Leu	tca Ser	2129

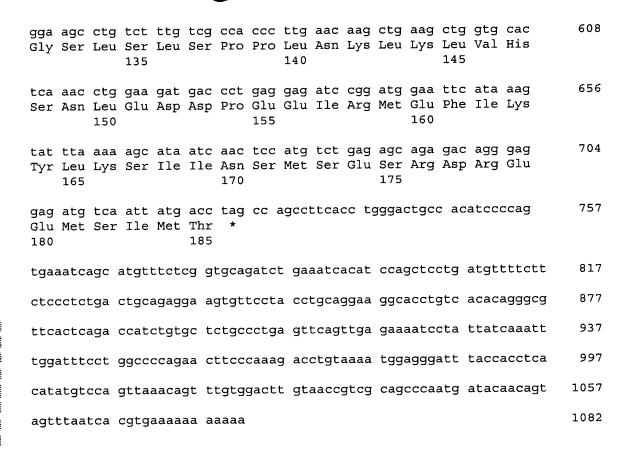
435 425 gac cga cat gca ttt gag aca gcg cgg cgt ctg agc cat gat aac atg 2177 Asp Arg His Ala Phe Glu Thr Ala Arg Arg Leu Ser His Asp Asn Met 445 450 gag gag aag gtg ccc tgg gtg tgc ccc cgg ccc ctg gtt ctg ccc tca 2225 Glu Glu Lys Val Pro Trp Val Cys Pro Arg Pro Leu Val Leu Pro Ser 465 2273 cct ctt qtc acc cct qqa agc aat agt cag gag cga tat atc cag gct Pro Leu Val Thr Pro Gly Ser Asn Ser Gln Glu Arg Tyr Ile Gln Ala 480 475 gag cgg gag aag gga atc ctt cag gag ctc ttc ctg aac aag gag agt 2321 Glu Arg Glu Lys Gly Ile Leu Gln Glu Leu Phe Leu Asn Lys Glu Ser 490 2369 cct cat gag cct gat cct gag ccc tac gag ccc ata ccc cct aaa ctc Pro His Glu Pro Asp Pro Glu Pro Tyr Glu Pro Ile Pro Pro Lys Leu 505 510 atc ccc cta gat gag gag tgt tcc atg gat gag act ccg tat gtt gag 2417 Ile Pro Leu Asp Glu Glu Cys Ser Met Asp Glu Thr Pro Tyr Val Glu act ctg gaa cct ggg ggg tca ggt ggc tca cct gat ggg gca gga ggc 2465 Thr Leu Glu Pro Gly Gly Ser Gly Gly Ser Pro Asp Gly Ala Gly Gly tcc aag ttg cct cca gtt ctg gcc aat ctt atg gga agc atg ggt gct 2513 Ser Lys Leu Pro Pro Val Leu Ala Asn Leu Met Gly Ser Met Gly Ala gga aag ggc ccc caa ggc cct gga gga ggc att aat gtc caa gag 2561 Gly Lys Gly Pro Gln Gly Pro Gly Gly Gly Ile Asn Val Gln Glu 575 atc ctc acc tcc atc atg ggt agc cca aac agt cat cct tca gag gaa 2609 Ile Leu Thr Ser Ile Met Gly Ser Pro Asn Ser His Pro Ser Glu Glu 595 585 590 cta ctg aaa caa cca gac tat tcg gac aag atc aag cag atg ctg gtg 2657 Leu Leu Lys Gln Pro Asp Tyr Ser Asp Lys Ile Lys Gln Met Leu Val 615 600 605 cca cat gga ctc cta ggc cct ggc cca ata gcc aat ggt ttc cca cca 2705 Pro His Gly Leu Leu Gly Pro Gly Pro Ile Ala Asn Gly Phe Pro Pro 620 630 ggg ggt cct ggg ggc ccc aag ggc atg cag cac ttt ccc cct gga cct 2753 Gly Gly Pro Gly Gly Pro Lys Gly Met Gln His Phe Pro Pro Gly Pro 635 640 ggg gga cct atg cca ggt ccc cat gga ggc cct ggt ggg cca gtg ggt 2801 Gly Gly Pro Met Pro Gly Pro His Gly Gly Pro Gly Gly Pro Val Gly

cca Pro	cgt Arg 665	ctt Leu	ctg Leu	ggt Gly	cct Pro	cca Pro 670	ccc Pro	cct Pro	ccc Pro	cgg Arg	gga Gly 675	ggt Gly	gat Asp	ccc Pro	ttc Phe	2	849
													cgg Arg			2	897
													ggc Gly			2	945
gga Gly	aac Asn	gaa Glu	cct Pro 715	cct Pro	cct Pro	cct Pro	cct Pro	cct Pro 720	cca Pro	ttc Phe	cga Arg	ggc Gly	gcc Ala 725	aga Arg	gga Gly	2	993
ggt Gly	cgc Arg	tct Ser 730	gga Gly	gga Gly	gga Gly	ccc Pro	cca Pro 735	aat Asn	gga Gly	cga Arg	ggg Gly	ggc Gly 740	cct Pro	ggt Gly	gly aaa	3	3041
ggc Gly	atg Met 745	gtt Val	gga Gly	ggt Gly	ggt Gly	999 Gly 750	cat His	cgt Arg	cct Pro	cac His	gaa Glu 755	ggc Gly	cct Pro	ggt Gly	gly aaa	3	1089
ggc Gly 760	atg Met	ggc Gly	aac Asn	agc Ser	agt Ser 765	gga Gly	cat His	cgt Arg	ccc Pro	cac His 770	gaa Glu	ggc Gly	cct Pro	ggc Gly	ggt Gly 775	3	3137
ggc Gly	atg Met	gga Gly	agt Ser	999 Gly 780	cat His	cgc Arg	ccc Pro	cat His	gaa Glu 785	ggc Gly	cct Pro	ggt Gly	ggt Gly	agc Ser 790	atg Met	3	3185
ggt Gly	ggg Gly	ggt Gly	gga Gly 795	gga Gly	cat His	cgt Arg	ccc Pro	cac His 800	gaa Glu	ggc Gly	cct Pro	ggc Gly	ggt Gly 805	ggc Gly	atc Ile	3	3233
agt Ser	ggt Gly	ggc Gly 810	agt Ser	ggc Gly	cat His	cgt Arg	ccc Pro 815	cat His	gaa Glu	ggc Gly	cct Pro	ggc Gly 820	gga Gly	gga Gly	atg Met	3	3281
ggt Gly	gcc Ala 825	ggt Gly	ggt Gly	gga Gly	cat His	cgc Arg 830	ccc Pro	cac His	gaa Glu	ggc Gly	cct Pro 835	ggc Gly	gga Gly	agc Ser	atg Met	3	3329
ggt Gly 840	gga Gly	agt Ser	ggt Gly	gga Gly	cat His 845	cgt Arg	ccc Pro	cat His	gaa Glu	ggc Gly 850	cct Pro	gga Gly	cac His	GJA aaa	999 Gly 855	3	3377
ccc Pro	cat His	ggc Gly	cac His	cgg Arg 860	cct Pro	cat His	gat Asp	gtc Val	cct Pro 865	ggt Gly	cac His	cga Arg	ggc Gly	cat His 870	gac Asp	3	3425
													ggt Gly 885			3	3473

cac ggg gga His Gly Gly 890	ggg ggc cac Gly Gly His	cga ggg ca Arg Gly H: 895	ac gat gga is Asp Gly	ggc cac agc Gly His Ser 900	cat gga His Gly	3521
gga gac atg Gly Asp Met 905	tca aac cgc Ser Asn Arg	cct gtc to Pro Val C	gc cga cat ys Arg His	ttc atg atg Phe Met Met 915	aag ggc Lys Gly	3569
aac tgc cgc Asn Cys Arg 920	tat gag aac Tyr Glu Asn 925	aac tgt g Asn Cys A	cc ttc tac la Phe Tyr 930	cac ccg ggt His Pro Gly	gtc aat Val Asn 935	3617
ggg ccc ccc Gly Pro Pro	ctg ccc tag Leu Pro * 940	ggacca tt	tgcctgcc ct	gttcacac aa	ccctgtg	3671
gactgcagcc	tegetettte e	accctgtta	tggcttctgt	gaggcccatt	ttcccttttc	3731
cccagctgat	gaggagccgg c	cccctcagt	tcccacttgc	ttgggttcct	gggggttttc	3791
tgatcactgg	tgcgcattga t	gtacatatt	ttcctccagt	ctggggagga	gagagactgg	3851
aaacgttcct	ggactgctga a	gaggagacc	cagttggctt	cactttttga	gaagattcgc	3911
cctgtacccc	aaaccccttt c	cagtattac	ccttaatgct	tgagaaccta	aagctggtta	3971
tcctggcgaa	cacccctacc c	ttctattgc	gggtccccac	atgcacacag	aactctgaca	4031
caggatcagc	tgcacttaag a	aatcatccc	agctaagttc	attattcctc	atggggtggg	4091
gagatgctga	aaggggtatt g	tatatccca	ctgcactgag	agggctcaat	cagctggatt	4151
tgagttctgg	aacacacatc a	tccccaccc	ctccccagc	gtgggctcac	cattcttagt	4211
cctttctcaa	gtgggacctt c	aactttctg	tgaacaccca	gtctgcgtcc	tgggtctgct	4271
aggttcgatg	atggcgaact c	gtatctgca	tccggtgcaa	gttttagctg	gcagaggtga	4331
gaccggtggt	gctggtctgc c	tttgccaac	tatagccagt	ctggagactt	gataaaatac	4391
ttcagtgaga	ccagcttctc a	tcaacttgg	gcccggcgtg	ctgggcctga	aagtcacact	4451
acatgcactg	cctttgggag t	cagctcact	ccctgctccc	acctggaacc	ttgccagcgt	4511
gaaggaggct	tccaggtact t	caccctgtc	aaccacctct	gaatccccac	caggcgcctt	4571
cctgggtgga	ttcaacaaga t	gattttgcc	ctttcccagt	tctctccttc	actttggcat	4631
cagttgtttt	ctatgaaaac a	agtggattgg	ttgggttttg	tgcagggtct	tgggttagag	4691
ccaaaatgga	tttgaggatg a	agtattttt	tttttggttt	tgtatatttt	gtacattaat	4751
aataaacagt	ggaaagagaa a	aaaaaaaaa				4781

(210) 25	
<211> 968	
<212> DNA	
<213> Homo sapiens	
<220>	
<221> CDS	
<222> (184)(411)	
<220>	
<221> misc_feature	
<222> (1)(968)	
<223> n = a,t,c or g	
<400> 29	
ttttttttt attattatac tttaagtttt agggtacatg tgcacaacgt gcaggtttgt	60
tacatatgta tacatgtgcc atgttagtgt gctgcaccca ttaactcgtc atttagcatt	120
aggtatatet ectacageta tecetecece ettececeae eccacaaaag gteceagtgt	180
gtg atg ttc ccc ttc ctg tgt cca tgt gtt ctc att gtt caa ttc cca	228
Met Phe Pro Phe Leu Cys Pro Cys Val Leu Ile Val Gln Phe Pro	
1 5 10 15	
the second secon	276
cct atg agt gag aac atg tgg gtt tgg ttt ttt tgt cct tgc gat agt	270
Pro Met Ser Glu Asn Met Trp Val Trp Phe Phe Cys Pro Cys Asp Ser	
20 25 30	
ttg ctg aga atg atg gtt tcc agc ttc atc cat gtc cct gcg aag gac	324
Leu Leu Arg Met Met Val Ser Ser Phe Ile His Val Pro Ala Lys Asp	
35 40 45	
atg aac tca ccc ttt ttt atg gaa tac tac aca gcc ata aaa agg aat	372
Met Asn Ser Pro Phe Phe Met Glu Tyr Tyr Thr Ala Ile Lys Arg Asn	
50 55 60	
and the set one of the second	421
gac aac aca tcc ctt gca ggg aca tgg atg gag caa tag gccattatcc	120
ASP ASI THE SET LEG ATO CTY THE TEP MES CER CER	
65 70 75	
gtagcaaact aatgcaggaa gagaaaaccc agtactgcat gttctcactt ataagtggga	481
gctaaatgat gagaacacat gaacacaaag aggggaacag acactagggc cgtttagaag	541
ttggcgggtg gtttgctttt ttntttagnt acangattta ttagnaatgg gtactaggct	601
gaataccgtt gtggatggta gtaatcgggt gaacaaagcg cccatgttca caattttagc	663
ttattttttc tggttttcgt catggtaccc cttgtgagtg tgagttatat gtttggttga	72
tatttatcgg ttattggatt tggtccctgg gtggtgtttt cgtggagtgg tggggttgtt	783
gtgattettg tgtttggetg tggtgtttaa aagaggtgta ggttggtgac eegggtgttg	841
tccacqttat tqqctttqat qaqqccccct tttgtgggtg gtgtttggtg ttgttttggt	90:

	ctgg gtgt		jtg <u>s</u>	geeet	attt	g tg	gcttt	ttgt	tgg	gtagt	.ggc	ttco	tggt	gg t	cggt	gagcg	961 968
		<21 <21 <21 <22 <22	20> 21> 0	DNA Homo	sapi												
	tttt)0> 3 aaa c		gatag	jc ag	gtaco	ccago	tgg	gctag	ıcgt	ttaa	actt	aa g	gcttg	ggtacc	60
,= <u>:</u>																cctgac	120
See News Treat State Hirty H. T.	gcg	cege	ete d	ecgto	gget	ic cg	ggcc	ggcta	a ago	eegeg	gcg	gaca	aact	Met		g aaa 1 Lys	176
Sur Verill Sur Verill	gcc Ala	aag Lys 5	atc Ile	ctc Leu	ttc Phe	gtg Val	ggg Gly 10	cct Pro	tgc Cys	gag Glu	agt Ser	gga Gly 15	aaa Lys	act Thr	gtt Val	ttg Leu	224
	gcc Ala 20	aac Asn	ttt Phe	ctg Leu	aca Thr	gaa Glu 25	tct Ser	tct Ser	gac Asp	atc Ile	act Thr 30	gaa Glu	tac Tyr	agc Ser	cca Pro	acc Thr 35	272
	caa Gln	gga Gly	gtg Val	agg Arg	atc Ile 40	cta Leu	gaa Glu	ttt Phe	gag Glu	aac Asn 45	ccg Pro	cat His	gtt Val	acc Thr	agc Ser 50	aac Asn	320
					ggc Gly												368
	gct Ala	aag Lys	ttt Phe 70	gag Glu	tcc Ser	tgc Cys	tgg Trp	ccg Pro 75	gcc Ala	ctg Leu	atg Met	aag Lys	gat Asp 80	gct Ala	cat His	gga Gly	416
					ttc Phe												464
	gag Glu 100	atg Met	tgg Trp	tat Tyr	tcc Ser	tgc Cys 105	ttt Phe	gtc Val	caa Gln	cag Gln	ccg Pro 110	tcc Ser	tta Leu	cag Gln	gac Asp	aca Thr 115	512
					att Ile 120												560

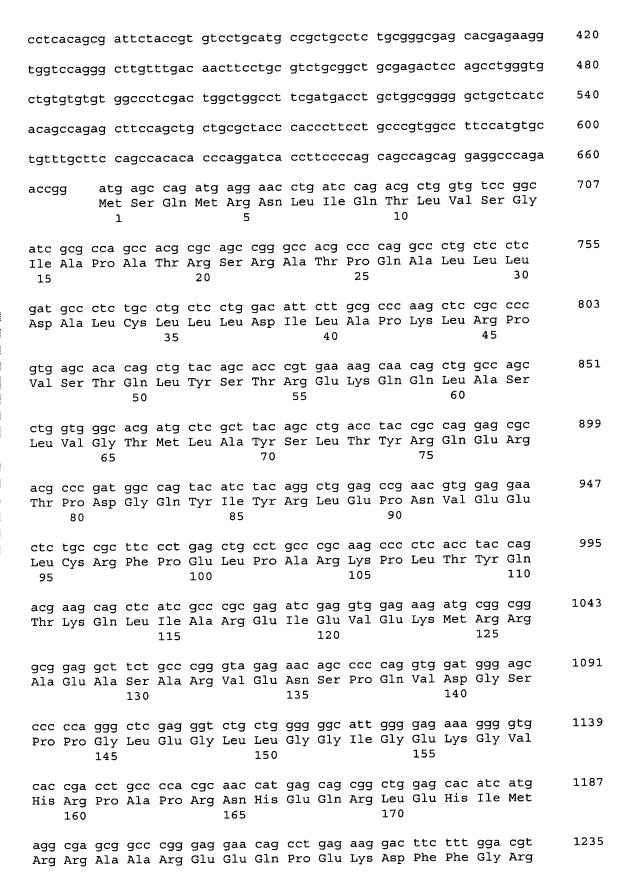


<211> 1517 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (666)..(1406) <220> <221> misc_feature <222> (1)...(1517)

<223> n = a,t,c or g

<210> 31

ctcgaaatcg atactttgcc ggaccggnnc ggnnnnccgg ggtcgacggg aggcaggagg 60 gccgacccag gggtgctggc cgcctctgt gagaaaactg acaatgacat ccgggcctgc 120 atcaacaccc tgcagttcct gtacagccgg ggccagcggg agctgagcgt gcgggacgtg 180 caggccacac gcgtgggct caaggaccag cgcagagggc tcttctcggt gtggcaggag 240 gtcttccagc tgctcgagc ccagagcacc ccacctgcag gcgccgtgt ggccaggacc 300 ccgcctgcc tgctgacaca ctcctgctgg gtgacggga cgcggggtcc ctcacctccg 360



185 190 180 175 1283 gtg gtc gtc agg agc aca gca gtc ccg agt gca ggg gac acg gcc ccg Val Val Val Arg Ser Thr Ala Val Pro Ser Ala Gly Asp Thr Ala Pro 200 195 gag cag gac tca gtg gag cgg cgc atg ggc aca gcg gtg ggc agg agc 1331 Glu Gln Asp Ser Val Glu Arg Arg Met Gly Thr Ala Val Gly Arg Ser 215 210 1379 gag gtc tgg ttc cgc ttc aac gag ggt gtc tcc aac gcc gtg cgg cgc Glu Val Trp Phe Arg Phe Asn Glu Gly Val Ser Asn Ala Val Arg Arg 225 230 age etg tae ate agg gae ttg ete tag ttete tgageegegg acatgeeete 1431 Ser Leu Tyr Ile Arg Asp Leu Leu * 245 240 gcattgcttc ccgcagagtg cagagacagg aagctggaga tgtctttata aagtcacacc 1491 1517 tttacagact gtaaaaaaaa aaaaaa <210> 32 <211> 618 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (498)..(572) <400> 32 ttttgcctaa gatacaatga taagtaaaaa catgttacag agtaataacg tgttgcatag 60 gacaatactg taaatgtggt tcattccatc aacagacact gagccctgcg acgtgcctgg 120 ctctattcta catctgaggg acacaaggtg aacaagacca ggccactgta ttcaacatct 180 acatttaatg gaaatttttg aaagaagact tgagagatta taacagtggt cttcagcatc 240 agggagtagg cctagaagaa gaggaggtca agaagtggct tttcttatta tcttcttaac 300 tottcaaatt tttactatga gcaattatta ttttttatta aaattttagg ccaggottat 360 ggctgtaatc ctagcagttt gggaggccaa ggtgagcgga tcacttgagg ttgggagttc 420 gagaccagcc tgaccaacat ggagaaactc tgtctctact taaaaaaaaa tacaaaatta 480 atg cct gta gtc cca gct act cag gag act gag 530 gccaggcatg gtggcac Met Pro Val Val Pro Ala Thr Gln Glu Thr Glu 579 gca ggg gaa ttg cct gaa cct ggg aga cag agg ttg caa tga gccaaga Ala Gly Glu Leu Pro Glu Pro Gly Arg Gln Arg Leu Gln *

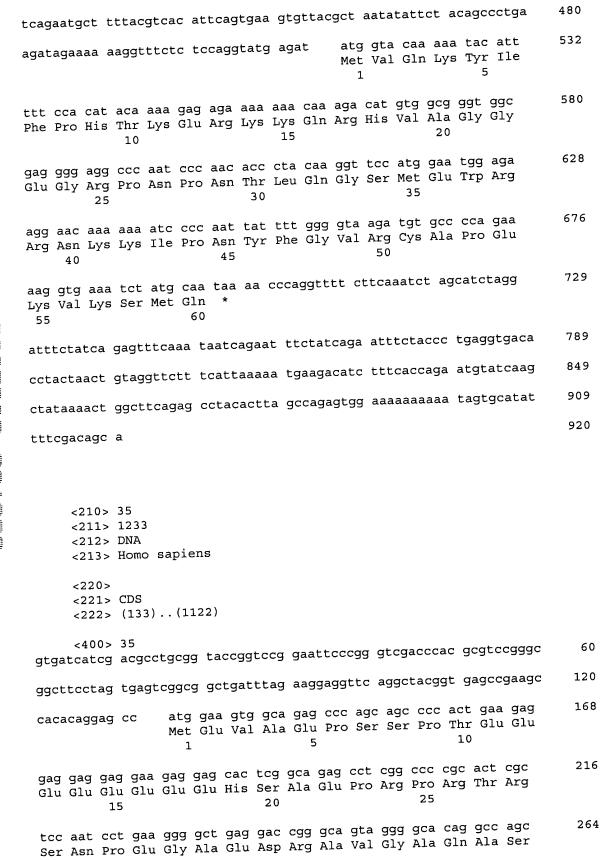
15 20 25

tcacgcctcg tgccgaattc ttggcctcga gggccaaat

<210> 33 <211> 1188 <212> DNA <213> Homo sapiens	
<220> <221> CDS <222> (313)(849)	
<400> 33 aaggateett aattaaatta ateeeeeee eeeggggaga aaegttetea etegetetet	60
gctcgctgcg ggcgctcccc gccctctgct gccagaacct tggggatgtg cctagacccg	120
gcgcagcaca cgtccgggcc aaccgcgagc agaacaaacc tttggcgggc ggccaggagg	180
ctccctccca gccaccgccc ccctccagcg cctttttttc cccccataca atacaagatc	240
ttccttcctc agttccctta aagcacagcc cagggaaacc tcctcacagt tttcatccag	300
ccacgggcca gc atg tct ggg ggc aaa tac gta gac tcg gag gga cat Met Ser Gly Gly Lys Tyr Val Asp Ser Glu Gly His 1 5 10	348
ctc tac acc gtt ccc atc cgg gaa cag ggc aac atc tac aag ccc aac Leu Tyr Thr Val Pro Ile Arg Glu Gln Gly Asn Ile Tyr Lys Pro Asn 15 20 25	396
aac aag gcc atg gca gac gag ctg agc gag aag caa gtg tac gac gcg Asn Lys Ala Met Ala Asp Glu Leu Ser Glu Lys Gln Val Tyr Asp Ala 30 35 40	444
cac acc aag gag atc gac ctg gtc aac cgc gac cct aaa cac ctc aac His Thr Lys Glu Ile Asp Leu Val Asn Arg Asp Pro Lys His Leu Asn 45 50 55 60	492
gat gac gtg gtc aag att gac ttt gaa gat gtg att gca gaa cca gaa Asp Asp Val Val Lys Ile Asp Phe Glu Asp Val Ile Ala Glu Pro Glu 65 70 75	540
ggg aca cac agt ttt gac ggc att tgg aag gcc agc ttc acc acc ttc Gly Thr His Ser Phe Asp Gly Ile Trp Lys Ala Ser Phe Thr Thr Phe 80 85 90	588
act gtg acg aaa tac tgg ttt tac cgc ttg ctg tct gcc ctc ttt ggc Thr Val Thr Lys Tyr Trp Phe Tyr Arg Leu Leu Ser Ala Leu Phe Gly 95 100 105	636
atc ccg atg gca ctc atc tgg ggc att tac ttc gcc att ctc tct ttc Ile Pro Met Ala Leu Ile Trp Gly Ile Tyr Phe Ala Ile Leu Ser Phe	684

	=	i i
:	1	į
:	7	teen7
:	they they	1
	4,11411	il.
•	Ä	Time.
:	the Hand	1111
:	į	'lent
	į	
;	=	i i
:	li con	Ī
	=	
	Harrie	ì
:	144	-
***	3	1
****	=======================================	Speed Street

110 ctg cac atc tgg gca gtt gta cca tgc att aag agc ttc ctg att gag 732 Leu His Ile Trp Ala Val Val Pro Cys Ile Lys Ser Phe Leu Ile Glu 130 125 att cag tgc atc agc cgt gtc tat tcc atc tac gtc cac acc gtc tgt 780 Ile Gln Cys Ile Ser Arg Val Tyr Ser Ile Tyr Val His Thr Val Cys 150 gac cca ctc ttt gaa gct gtt ggg aaa ata ttc agc aat gtc cgc atc 828 Asp Pro Leu Phe Glu Ala Val Gly Lys Ile Phe Ser Asn Val Arg Ile 170 165 160 aac ttg cag aaa gaa ata taa at gacatttcaa ggatagaagt atacctgatt 881 Asn Leu Gln Lys Glu Ile * 175 ttttttcctt ttaattttcc tggtgccaat ttcaagttcc aagttgctaa tacagcaaca 941 atttatgaat tgaattatct tggttgaaaa taaaaagatc actttctcag ttttcataag 1001 tattatgtct cttctgagct atttcatcta tttttggcag tctgaatttt taaaacccat 1061 ttaaattttt ttccttacct ttttatttgc atgtggatca accatcgctt tattggctga 1121 gatatgaaca tattgttgaa aggtaatttg agagaaatat gaagaactga ggaggaaaaa 1181 1188 aaaaaaa <210> 34 <211> 920 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (515)..(697) <400> 34 taagcttgcg gccgcaattt ttttttttt ttttttgtat tttttggtag agacgggatt 60 tcactatgtt ggtcaggctg gtctcgaact cccgaccgca agtgatccac ccgccttggc 120 ctcccaaagt gctgggatta caagcttgag ccactgcacc cagcctggaa agtatattta 180 tgaaaggttt gcactccaca aaagcatctt tgctagggtg tcaaggaaga gatcactaaa 240 ccaaccccaa cacatccata caattccagc aatctagaga gggctggtcc ttttcctttt 300 ctggattatt ttctgttctc agtaaaacaa gtatttactg tgatactgaa acactgggaa 360 attaacactg attaagatat tttaaacact gagtcttaat tataacagaa ccagttttca 420



40 30 gtg ggc agc cgc agc gag ggt gag ggt gag gcc gcc agt gct gat gat 312 Val Gly Ser Arg Ser Glu Gly Glu Gly Glu Ala Ala Ser Ala Asp Asp 50 45 ggg agc ctc aac act tca gga gcc ggc cct aag tcc tgg cag gtg ccc 360 Gly Ser Leu Asn Thr Ser Gly Ala Gly Pro Lys Ser Trp Gln Val Pro ccg cca gcc cct gag gtc caa att cgg aca cca agg gtc aac tgt cca 408 Pro Pro Ala Pro Glu Val Gln Ile Arg Thr Pro Arg Val Asn Cys Pro 85 80 gag aaa gtg att atc tgc ctg gac ctg tca gag gaa atg tca ctg cca 456 Glu Lys Val Ile Ile Cys Leu Asp Leu Ser Glu Glu Met Ser Leu Pro 95 aag ctg gag tcg ttc aac ggc tcc aaa acc aac gcc ctc aat gtc tcc 504 Lys Leu Glu Ser Phe Asn Gly Ser Lys Thr Asn Ala Leu Asn Val Ser cag aag atg att gag atg ttc gtg cgg aca aaa cac aag atc gac aaa 552 Gln Lys Met Ile Glu Met Phe Val Arg Thr Lys His Lys Ile Asp Lys 130 agc cac gag ttt gca ctg gtg gtg gtg aac gat gac acg gcc tgg ctg 600 Ser His Glu Phe Ala Leu Val Val Val Asn Asp Asp Thr Ala Trp Leu tet gge etg ace tee gae eec ege gag etc tgt age tge etc tat gat 648 Ser Gly Leu Thr Ser Asp Pro Arg Glu Leu Cys Ser Cys Leu Tyr Asp 160 ctg gag acg gcc tcc tgt tcc acc ttc aat ctg gaa gga ctt ttc agc 696 Leu Glu Thr Ala Ser Cys Ser Thr Phe Asn Leu Glu Gly Leu Phe Ser 180 175 ctc atc cag cag aaa act gag ctt ccg gtc aca gag aac gtg cag acg 744 Leu Ile Gln Gln Lys Thr Glu Leu Pro Val Thr Glu Asn Val Gln Thr 195 190 att ccc ccg cca tat gtg gtc cgc acc atc ctt gtc tac agc cgt cca 792 Ile Pro Pro Pro Tyr Val Val Arg Thr Ile Leu Val Tyr Ser Arg Pro 215 210 205 cct tgc cag ccc cag ttc tcc ttg acg gag ccc atg aag aaa atg ttc 840 Pro Cys Gln Pro Gln Phe Ser Leu Thr Glu Pro Met Lys Lys Met Phe 230 225 cag tgc cca tat ttc ttc ttt gac gtt gtt tac atc cac aat ggc act 888 Gln Cys Pro Tyr Phe Phe Asp Val Val Tyr Ile His Asn Gly Thr 240 gag gag aag gag gag atg agt tgg aag gat atg ttt gcc ttc atg 936 Glu Glu Lys Glu Glu Glu Met Ser Trp Lys Asp Met Phe Ala Phe Met 255

